## Richard M. Kostrzewa *Editor*

# Handbook of Neurotoxicity

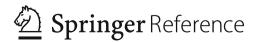


Handbook of Neurotoxicity

Richard M. Kostrzewa Editor

## Handbook of Neurotoxicity

With 244 Figures and 38 Tables



*Editor* Richard M. Kostrzewa Department of Biomedical Sciences Quillen College of Medicine East Tennessee State University Johnson City, TN, USA

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#### Preface

*Handbook of Neurotoxicity* is a compendium intended to provide an overview on neurotoxins. In addition, processes and mechanisms attending neuronal necrosis, necroptosis, and apoptosis are discussed in detail. Also, glial perturbations, as these relate to associated neuronal dysfunction, and both in vivo and in vitro methodological approaches toward the study and use of neurotoxins are discussed.

The *first theme* in the *Handbook* on selective neurotoxins provides a listing of individual neurotoxins, with a description of relative selectivity of each, including cellular mechanisms of action and overall neurologic and behavioral outcome deriving from these neurotoxins' effects. The *second section*, on dopamine-derived neurotoxins, provides highlights of neuronal mechanisms associated with the production of reactive dopamine quinones and semiquinones and reactive oxygen/nitrogen species formed by the reactive dopamine analogs. These processes and mechanisms are related to purported genesis of Parkinson's disease.

The *third section*, on glutamate receptor agonists, is focused on the theme of neuronal excitotoxicity as an outcome of over-excitation of neurons possessing kainate receptors, AMPA (alpha-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid) receptors, and NMDA (N-methyl-D-aspartate) receptors. These excitotoxins are discussed in detail as to their known and putative roles in neurodegenerative and psychiatric disorders. Similarly, the roles of exogenous and endogenous glutamate receptor antagonists as neuroprotectants and as neurotoxins per se are described.

The *fourth section* highlights the roles of neurotrophins in neuronal development and relates their involvement in a host of neurotoxicity events in neurologic and neurodegenerative disorders. The *last section* of the *Handbook*, focused on diseases and disorders relevant to neurotoxins, describes processes and mechanisms associated with specific neural disorders, including effects of some of the most prominent neurotoxins in relation to human dysfunction. Additionally, impugned neurotoxicity mechanisms in neurological disorders, protective strategies for averting and delaying onset of neurologic and psychiatric disorders, and highlights of newer strategies towards therapy are also discussed.

*Handbook of Neurotoxicity* is intended to be useful to students, scientists, and clinicians as an initial introduction to neurotoxin-associated processes. Simultaneously, the *Handbook* is expected to be a valuable guide for those employing specific neurotoxins as there are specific details on the use of individual neurotoxins along with dosages, delineation of mechanisms of action, expected neuronal

degenerative effects, behavioral outcome, and possible duration of effect and recovery from damage.

In essence, the *Handbook* is to represent an authoritative reference resource on neurotoxins and neurotoxicity processes and mechanisms in the medical and scientific arena.

#### **Editor-in-Chief**

Richard M. Kostrzewa, Ph.D., Dr.h.c., obtained a Ph.D. (1971) in Pharmacology from the University of Pennsylvania. After positions as Research Pharmacologist at the Veterans Administration Medical Center in New Orleans, Louisiana (1971–1975); adjunct Assistant Professor in the Department of Pharmacology at Tulane Medical School (1974-1976) and Department of Psychology at the University of New Orleans (1974–1977); and Associate Professor at the Louisiana University Medical Center in New Orleans (1975–1978), Dr. Kostrzewa joined the faculty of Pharmacology at the newly established Ouillen College of Medicine at East Tennessee State University in Johnson City, Tennessee (Assoc. Prof./ Professor, 1978 to present). He actively collaborated in research for 20 years with Prof. Ryszard Brus of the Medical University of Silesia in Katowice, Poland, where Dr. Kostrzewa was Visiting Professor (1997–2003) and from which he received the Doctorate Honoris Causa (2005). In 2001, Dr. Kostrzewa was made an Honorary Member of the Polish Pharmacological Society. His research focus has remained on selective neurotoxins, with applications in the areas of nerve regeneration, neurodegenerative disorders, and dopamine receptor supersensitivity. He has authored and edited several books and more than 200 scientific papers, including the article coauthored with Dr. David M. Jacobowitz, his former Dissertation Advisor, and designated as a "Citation Classic" in Current Contents: "Pharmacological Actions of 6-Hydroxydopamine." Dr. Kostrzewa was the inaugural President of the Neurotoxicity Society and served as the Society's Treasurer for ten years. He has been Editorin-Chief of the journal *Neurotoxicity Research* from the time of its inception (1998).

#### **Section Editors**

**Trevor Archer** obtained an Honour's degree in Psychology and Physiology from Newcastle-upon-Tyne University in 1974, and a Ph.D. from Uppsala University, Uppsala, Sweden, in 1979. From 1978 to 1988, he was Head of CNS Drugs at Astra Läkemedel AB. He was then appointed Professor and Chair of Biological Psychology at the University of Gothenburg, Sweden. He has been Guest or Visiting Professor at: the University of Hawaii at Manoa; Örebro University; Karlstad University; Kalmar University; and Madrid University. He received an Outstanding Lifetime Achievement Award from St. Joseph's College, North Point in 2010, and Honorary Doctorship, Professorship, and Membership Awards from the Polish Academy of Science Pharmacological Society in 2013.

**Gilles Guillemin** is CORE Professor and Co-Director of the newly-created "MND and Neurodegenerative Diseases Research Centre" at Macquarie University. He is an Honorary Associate of the University of Sydney and the Visiting Senior Research Fellow at the Australasian Research Institute, Sydney Adventist Hospital, Wahroonga. Professor Guillemin has published over 110 papers and edited a book, *Neuroanatomie fonctionnelle: La cellule microgliale.* He has also contributed to 12 other books and presented at over 90 conferences. Dr. Guillemin is heavily involved in tryptophan research, as his group is one of the world's leading research groups working on neuroactive metabolites derived from tryptophan. He is the newly-elected President of both the International Society for Tryptophan Research (ISTRY) and the Neurotoxicity Society (NTS). His current research interest is studying the involvement of tryptophan catabolism (via the kynurenine pathway) in human neurodegenerative diseases such as amyotrophic lateral sclerosis, multiple sclerosis, Parkinson's disease, Alzheimer's disease, and brain tumors.

**Juan Segura-Aguilar** studied chemistry with special mention in microbiology at the University of Stockholm, Sweden, and received his bachelor's degree in 1983. In 1989, he received his Ph.D. in biochemistry from the University of Stockholm. In 1999, he was awarded a stay at the University of Valencia by the European Community. From 1993 to 1998, he was at Uppsala University, Sweden, where he obtained "Docent" competence in 1994. In 1998, he moved to Chile and has been Full Professor at the University of Chile. He has been awarded several competitive research grants, including FONDECYT grants. Currently, he leads a research group

that studies the role of dopamine oxidation in the degenerative process of the nigrostriatal dopaminergic system. With Prof. Richard Kostrzewa, he founded and organized the Neurotoxicity Society, and has also organized several international NTS meetings, both regular and satellite.

**Xin-Fu Zhou** received his undergraduate medical degree from Third Military Medical University in China in 1977, M.Sc. in neurology from Beijing PLA Postgraduate College in 1982, and Ph.D. in biochemistry from the University of Melbourne in 1990. He was awarded an Australian Postdoctoral Fellowship by the Australian Research Council (ARC) in 1991 and completed his postdoctoral research at Flinders University. Since 1997, he has been awarded four times Research Fellowships and named Senior Research Fellow by the National Health and Medical Research Council (NHMRC). He has been awarded numerous competitive research grants by ARC and NHMRC. His main research interest has been in neurological disorders. Recently, he is interested in functions of proneurotrophins and their receptors in neurological diseases such as Alzheimer's disease, spinal cord injury, and depression. Currently, he is NHMRC Senior Research Fellow and Research Chair in Neuroscience at University of South Australia.

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#### Contributors

Seray Adams School of Medical Sciences, Faculty of Medicine, University of New South Wales, Sydney, Australia

**Jonas I. Addae** Department of Preclinical Sciences, University of the West Indies, St Augustine, Trinidad and Tobago

**Pabla Aguirre** Department of Biology, Faculty of Sciences, University of Chile, Casilla, Santiago, Chile

**Ulises Ahumada-Castro** Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

**Ramiro D. Almeida** Center for Neuroscience and Cell Biology (CNC), University of Coimbra, Coimbra, Portugal

**Stefanie C. Altieri** David Geffen School of Medicine, Department of Chemistry and Biochemistry, Semel Institute for Neuroscience and Human Behavior, and Hatos Center for Neuropharmacology, University of California, Los Angeles, CA, USA

Anne M. Andrews David Geffen School of Medicine, Department of Chemistry and Biochemistry, Semel Institute for Neuroscience and Human Behavior, and Hatos Center for Neuropharmacology, University of California, Los Angeles, CA, USA

**Trevor Archer** Department of Psychology, University of Gothenburg, Gothenburg, Sweden

**Sara Ares-Santos** Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC) and CIBERNED, ISCIII, Madrid, Spain

Michael Aschner Department of Pediatrics, Vanderbilt University, Nashville, TN, USA

Department of Molecular Pharmacology, Albert Einstein College of Medicine, Bronx, NY, USA

Valeriya Avdoshina Department of Neuroscience, Georgetown University Medical Center, Washington, DC, USA

Daiana Silva Ávila Universidade Federal do Pampa, Uruguaiana, RS, Brazil

Alessia Bachis Department of Neuroscience, Georgetown University Medical Center, Washington, DC, USA

**Carlos Barcia** Clinical and Experimental Neuroscience / Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (NiCE–CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

**Eugenio Barone** Institute of Pharmacology, Catholic University School of Medicine, Rome, Italy

Department of Chemistry, Center of Membrane Sciences, and Sanders-Brown Center on Aging, University of Kentucky, Lexington, KY, USA

Shannon Beasley UQ Centre for Clinical Research, The University of Queensland, Brisbane, QLD, Australia

Simon A. Bedard International Collaboration on Repair Discoveries, The University of British Columbia, Vancouver, BC, Canada

**Verónica Bisagno** Instituto de Investigaciones Farmacológicas (ININFA-UBA-CONICET), Buenos Aires, Argentina

Aleksandra Bortel Departments of Neurology, Neurosurgery, and Biomedical Engineering, Montreal Neurological Institute, McGill University, Montreal, QC, Canada

**Mariza Bortolanza** Departamento MFP-Fisiologia, Faculdade de Odontologia de Ribeirão Preto, Universidade de São Paulo (USP), Center for Interdisciplinary Research on Applied Neurosciences (NAPNA), Ribeirão Preto, SP, Brazil

Luisa Bracci-Laudiero Institute of Translational Pharmacology, National Research Council of Italy (CNR), Rome, Italy

**Nady Braidy** Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

**José M. Bravo-San Pedro** Departamento de Bioquímica y Biología Molecular y Genética, Facultad de Enfermería y TO, Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas (CIBERNED), Universidad de Extremadura, Cáceres, Spain

George R. Breese Department of Psychiatry and Pharmacology, Bowles Center for Alcohol Studies, UNC School of Medicine, Chapel Hill, NC, USA

Amanda J. Bressler David Geffen School of Medicine, Department of Chemistry and Biochemistry, Semel Institute for Neuroscience and Human Behavior, and Hatos Center for Neuropharmacology, University of California, Los Angeles, CA, USA **Bruce J. Brew** St. Vincent's Centre for Applied Medical Research, Darlinghurst, NSW, Australia

University of New South Wales, Sydney, Australia

Departments of Neurology and HIV Medicine, St. Vincent's Hospital, Sydney, NSW, Australia

**Ryszard Brus** Department of Nursing, High School of Strategic Planning, Dabrowa Górnicza, Poland

Bogusława Budziszewska Department of Toxicology, Jagiellonian University Medical College, Kraków, Poland

Institute of Pharmacology, Polish Academy of Sciences and Center of Excellence in Neuropsychopharmacology, Kraków, Poland

**D. Bustamante** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**D. Allan Butterfield** Department of Chemistry, Center of Membrane Sciences, and Sanders-Brown Center on Aging, University of Kentucky, Lexington, KY, USA

**Jean Lud Cadet** NIDA Intramural Program, Molecular Neuropsychiatry Research Branch, NIDA/NIH/DHHS, Baltimore, MD, USA

Shuowei Cai National Botulinum Research Center, and Department of Chemistry and Biochemistry, University of Massachusetts Dartmouth, Dartmouth, MA, USA

Elena Caldarazzo-Ienco Clinical Neurology, University of Pisa, Pisa, Italy

**C. M. Campuzano** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain

Jason R. Cannon School of Health Sciences, Purdue University, West Lafayette, IN, USA

Anna R. Carta Department of Biomedical Sciences, University of Cagliari, Cagliari, Italy

Rudy J. Castellani Division of Neuropathology, University of Maryland, Baltimore, MD, USA

**Pablo Caviedes** Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

**Raúl Caviedes** Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

**Jing Chen** Department of Neurology, The Second Affiliated Hospital of Soochow University, Suzhou City, Jiangsu, China

The Institute of Neuroscience, Soochow University, Suzhou City, Jiangsu, China

**Zhe-Yu Chen** Department of Neurobiology, School of Medicine, Shandong University, Jinan, Shandong, China

**Iwona Chmiel-Perzyńska** Laboratory of Cellular and Molecular Pharmacology, Department of Experimental and Clinical Pharmacology, Medical University of Lublin, Lublin, Poland

Soyoung Choi Department of Biological Sciences, Rutgers, The State University of New Jersey, Newark, NJ, USA

Belinda Cruse Department of Neurology, St. Vincent's Hospital, Sydney, NSW, Australia

**Teresa Cunha-Oliveira** Center for Neuroscience and Cell Biology (CNC), University of Coimbra, Coimbra, Portugal

D. Cutuli I.R.C.C.S. Santa Lucia Foundation, Rome, Italy

**Ru-Ping Dai** Department of Anesthesia, The Second Xiang-Ya Hospital of Central South University, Changsha, Hunan Province, China

**Margaret I. Davis** Section on Synaptic Pharmacology, Laboratory for Integrative Neuroscience, National Institute on Alcohol Abuse and Alcoholism, National Institutes of Health, Bethesda, MD, USA

**P. De Bartolo** Department of Psychology, Sapienza University of Rome, Rome, Italy

I.R.C.C.S. Santa Lucia Foundation, Rome, Italy

Ana Carolina de Castro Issy Departamento MFP-Fisiologia, Faculdade de Odontologia de Ribeirão Preto, Universidade de São Paulo (USP), Center for Interdisciplinary Research on Applied Neurosciences (NAPNA), Ribeirão Preto, SP, Brazil

**Elaine Del Bel** Departamento MFP-Fisiologia, Faculdade de Odontologia de Ribeirão Preto, Universidade de São Paulo (USP), Center for Interdisciplinary Research on Applied Neurosciences (NAPNA), Ribeirão Preto, SP, Brazil

Marek Derkacz Department of Endocrinology, Medical University of Lublin, Lublin, Poland

**Peter R. Dodd** School of Chemistry and Molecular Biosciences, The University of Queensland, St. Lucia Campus, Brisbane, Australia

**Ana Paula Marreilha dos Santos** Department of Toxicology and Food Sciences, Faculty of Pharmacy, University of Lisbon, Lisbon, Portugal

**Bruno Lopes dos Santos** Department of Neurosciences and Behavior Sciences, Medical School of Ribeirão Preto, University of São Paulo, Ribeirão Preto, SP, Brazil

**Kay L. Double** The University of Sydney, Discipline of Biomedical Science, School of Medical Science, Sydney Medical School, Sydney, Australia

**Carlos B. Duarte** Center for Neuroscience and Cell Biology (CNC) and Department of Life Sciences, University of Coimbra, Coimbra, Portugal

Luc Dupuis INSERM, U1118, Faculté de Médecine, Mécanismes centraux et périphériques de la neurodégénérescence, Université de Strasbourg, Strasbourg, France

**D. Esmar** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**P. Espina-Marchant** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**M. M. Essa** Department of Food Science and Nutrition, College of Agriculture and Marine Sciences, Sultan Qaboos University, Muscat, Oman

**Margaret Fahnestock** Department of Psychiatry and Behavioural Neurosciences, McMaster University, Hamilton, ON, Canada

**E. Fernández-Villalba** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

**Małgorzata Filip** Department of Toxicology, Faculty of Pharmacy, Jagiellonian University, Kraków, Poland

Laboratory of Drug Addiction Pharmacology, Institute of Pharmacology Polish Academy of Sciences, Kraków, Poland

Vanderlei Folmer Universidade Federal do Pampa, Uruguaiana, RS, Brazil

Francesco Fornai Human Anatomy, University of Pisa, Pisa, Italy

Neuromed, I.R.C.C.S, Pozzilli, IS, Italy

**José María Frade** Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC), Madrid, Spain

**Lucia Frau** Department of Biomedical Sciences, University of Cagliari, Cagliari, Italy

Anders Fredriksson Department of Neuroscience, Psychiatry, Ulleråker, University Hospital, Uppsala, Sweden

Wilma J. Friedman Department of Biological Sciences, Rutgers, The State University of New Jersey, Newark, NJ, USA

**José M. Fuentes** Departamento de Bioquímica y Biología Molecular y Genética, Facultad de Enfermería y TO, Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas (CIBERNED), Universidad de Extremadura, Cáceres, Spain

**Ferenc Fülöp** Institute of Pharmaceutical Chemistry and Research Group of Stereochemistry, Hungarian Academy of Sciences, University of Szeged, Szeged, Hungary

Maria F. Galindo Unidad de Neuropsicofarmacología Traslacional, Complejo Hospitalario Universitario de Albacete, Albacete, Spain

Pharmaceutical Technology School of Pharmacy, University of Castilla-La Mancha, Albacete, Spain

**P. Gebicke-Haerter** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

Department of Psychopharmacology, Central Institute of Mental Health, Mannheim, Germany

Howard E. Gendelman Department of Pharmacology and Experimental Neuroscience, University of Nebraska Medical Center, NE, USA

**Manfred Gerlach** University of Würzburg, Department of Child and Adolescent Psychiatry, Psychosomatics and Psychotherapy, Laboratory for Clinical Neurobiology, Würzburg, Germany

**A. Gómez** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

**Rubén Gómez-Sánchez** Departamento de Bioquímica y Biología Molecular y Genética, Facultad de Enfermería y TO, Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas (CIBERNED), Universidad de Extremadura, Cáceres, Spain

**Rosa A. González-Polo** Departamento de Bioquímica y Biología Molecular y Genética, Facultad de Enfermería y TO, Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas (CIBERNED), Universidad de Extremadura, Cáceres, Spain

Chandan Goswami National Institute of Science Education and Research, Institute of Physics Campus, Bhubaneswar, Orissa, India

**Noelia Granado** Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC) and CIBERNED, ISCIII, Madrid, Spain

**Giorgia Greco** Analytical Research Group, Wissenschaftszentrum Weihenstephan, Technische Universität München, Freising, Germany

**J. Timothy Greenamyre** Pittsburgh Institute for Neurodegenerative Diseases, University of Pittsburgh, Pittsburgh, PA, USA

Gilles J. Guillemin St. Vincent's Centre for Applied Medical Research, St. Vincent's Hospital, Sydney, Australia

Neuropharmacology Group, MND and Neurodegenerative Diseases Research Centre, Macquarie University, North Ryde, NSW, Australia

**Dongwei Guo** Departments of Pharmaceutical Sciences, University of Nebraska Medical Center, Omaha, NE, USA

**Glenda Halliday** Neuroscience Research Australia and University of New South Wales, Sydney, NSW, Australia

**A. J. Hannan** The Florey Institute of Neuroscience and Mental Health, The University of Melbourne, Melbourne, Australia

**G. Jean Harry** Neurotoxicology Group, National Toxicology Program Laboratory, National Institute of Environmental Health Sciences, Research Triangle Park, NC, USA

**X. J. He** Department of Veterinary Pathology, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan

Elizabeth Head Department of Molecular and Biomedical Pharmacology, and Sanders-Brown Center on Aging, University of Kentucky, Lexington, KY, USA

**M. Hermoso** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**M. Herrera-Marschitz** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**M. T. Herrero** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain

David Howells Florey Neuroscience Institutes, Melbourne Brain Centre, Heidelberg, VIC, Australia

S. Hoyer Department of Pathology, University of Heidelberg, Heidelberg, Germany

Richard A. Hughes Department of Pharmacology, University of Melbourne, VIC, Australia

**Matthew R. Hynd** School of Chemistry and Molecular Biosciences, The University of Queensland, St. Lucia Campus, Brisbane, Australia

Chrysanthy Ikonomidou Department of Neurology, University of Wisconsin, Madison, WI, USA

**T. Ito** Department of Veterinary Pathology, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan

**Joseph Jankovic** Parkinson's Disease Center and Movement Disorders Clinic, Department of Neurology, Baylor College of Medicine, Houston, TX, USA

Tharusha Jayasena Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

**Poul Henning Jensen** Department of Biomedicine, Aarhus University, Aarhus C, Denmark

**Joaquín Jordán** Grupo de Neurofarmacología, Facultad de Medicina de Albacete, Universidad de Castilla-La Mancha, Albacete, Spain

**Ulla Karilampi** Department of Psychiatry/Psychosis, Sahlgrenska University Hospital, Gothenburg, Sweden

**M. Kaul** Infectious and Inflammatory Disease Center, Sanford-Burnham Medical Research Institute, La Jolla, CA, USA

Matthew C. Kiernan Bushell Professor of Neurology, Brain and Mind Research Institute, University of Sydney, Sydney, Australia

Neuroscience Research Australia, Randwick, Sydney, NSW, Australia

**Anna E. King** Wicking Dementia Research and Education Centre, University of Tasmania, Hobart, Tasmania, Australia

Péter Klivényi Department of Neurology, University of Szeged, Szeged, Hungary

John P. Kostrzewa Department of Biomedical Sciences, Quillen College of Medicine, East Tennessee State University, Johnson City, TN, USA

North Alabama ENT Associates, Huntsville, AL, USA

**Richard M. Kostrzewa** Department of Biomedical Sciences, Quillen College of Medicine, East Tennessee State University, Johnson City, TN, USA

Stjepana Kovac UCL Institute of Neurology, University College London, London, UK

Christine Lund Kragh Department of Biomedicine, Aarhus University, Aarhus C, Denmark

**C. Kraus** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

Weronika Krzyżanowska Department of Toxicology, Jagiellonian University Medical College, Kraków, Poland

Ashutosh Kumar National Institute of Science Education and Research, Institute of Physics Campus, Bhubaneswar, Orissa, India

**Raj Kumar** National Botulinum Research Center and Department of Chemistry and Biochemistry, University of Massachusetts Dartmouth, Dartmouth, MA, USA

Audrey P. Le Department of Biological Sciences, Rutgers, The State University of New Jersey, Newark, NJ, USA

Aven Lee UQ Centre for Clinical Research, The University of Queensland, Brisbane, QLD, Australia

**Zhijie Li** Cellular and Molecular Biology, Pediatric Oncology Branch, National Cancer Institute, National Institute of Health, Bethesda, MD, USA

Chai K. Lim Department of Pharmacology, School of Medical Sciences, University of New South Wales, Sydney, Australia

MND and Neurodegenerative Diseases Research Group, Macquarie University, Sydney, Australia

Seung T. Lim Department of Neuroscience, Georgetown University Medical Center, Washington, DC, USA

**Stuart A. Lipton** Del E. Webb Center for Neuroscience, Aging, and Stem Cell Research, Sanford-Burnham Medical Research Institute, La Jolla, CA, USA

Noelia López-Sánchez Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC), Madrid, Spain

C. T. Loy School of Public Health, The University of Sydney, Sydney, Australia

Huntington Disease Service, Westmead Hospital, Sydney, Australia

Jian-Jun Lu Division of Health Sciences, University of South Australia, Adelaide, SA, Australia

**Rakesh Kumar Majhi** National Institute of Science Education and Research, Institute of Physics Campus, Bhubaneswar, Orissa, India

Zsófia Majláth Department of Neurology, University of Szeged, Szeged, Hungary

Cesare Mancuso Institute of Pharmacology, Catholic University School of Medicine, Rome, Italy

**Luigi Manni** Institute of Translational Pharmacology, National Research Council of Italy (CNR), Rome, Italy

**Romesh Markus** Department of Neurology, St. Vincent's Hospital, Darlinghurst, Sydney, NSW, Australia

**Marta Martínez-Vicente** Neurodegenerative Diseases Research Group, Vall d'Hebron Research Institute–CIBERNED, Barcelona, Spain

Department of Biochemistry and Molecular Biology, Autonomous University of Barcelona (UAB), Barcelona, Spain

**Wakako Maruyama** Department of Cognitive Brain Science, National Research Center for Geriatrics and Gerontology, Obu Aichi, Japan

**JoEllyn M. McMillan** Department of Pharmacology and Experimental Neuroscience, University of Nebraska Medical Center, Omaha, NE, USA

**Christopher A. McPherson** Neurotoxicology Group, National Toxicology Program Laboratory, National Institute of Environmental Health Sciences, Research Triangle Park, NC, USA

**Raja Mehanna** Parkinson's Disease Center and Movement Disorders Clinic, Department of Neurology, Baylor College of Medicine, Houston, TX, USA

**Natalia Mena** Department of Biology, Faculty of Sciences, University of Chile, Casilla, Santiago, Chile

Stephanie Merchant College of Medicine, University of Central Florida, Orlando, FL, USA

**Italo Mocchetti** Department of Neuroscience, Georgetown University Medical Center, Washington, DC, USA

**P. Morales** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**Rosario Moratalla** Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC) and CIBERNED, ISCIII, Madrid, Spain

Micaela Morelli Department of Biomedical Sciences, University of Cagliari, Cagliari, Italy

CNR Institute of Neuroscience, Cagliari, Italy

**Renée Morris** Translational Neuroscience Facility, School of Medical Sciences, The University of New South Wales, Sydney, NSW, Australia

**Eugene V. Mosharov** Department of Neurology, Columbia University, New York, NY, USA

**V. Munoz** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**Dennis L. Murphy** Laboratory of Clinical Science, National Institute of Mental Health, National Institutes of Health, Bethesda, MD, USA

**Simon S. Murray** Department of Anatomy and Neuroscience, The University of Melbourne, Parkville, VIC, Australia

**H. Nakayama** Department of Veterinary Pathology, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan

**Jun Nakumura** Department of Psychiatry, School of Medicine, University of Occupational and Environmental Health, Yahatanishi-ku, Kitakyushu, Japan

Makoto Naoi Department of Neurosciences, Gifu International Institute of Biotechnology, Kakamigahara Gifu, Japan

**T. Neira-Peña** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**Ewa Niedzielska** Department of Toxicology, Faculty of Pharmacy, Jagiellonian University, Kraków, Poland

**Mireia Niso-Santano** Departamento de Bioquímica y Biología Molecular y Genética, Facultad de Enfermería y TO, Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas (CIBERNED), Universidad de Extremadura, Cáceres, Spain

Gabriel Nowak Department of Pharmacobiology, Jagiellonian University Medical College, Kraków, Poland

Institute of Pharmacology, Polish Academy of Sciences and Center of Excellence in Neuropsychopharmacology, Kraków, Poland

**Marco T. Núñez** Department of Biology, Faculty of Sciences, University of Chile, Casilla, Santiago, Chile

Victoria O'Collins Florey Neuroscience Institutes, Melbourne Brain Centre, Heidelberg, VIC, Australia

Jason B. Ochroch David Geffen School of Medicine, Department of Chemistry and Biochemistry, Semel Institute for Neuroscience and Human Behavior, and Hatos Center for Neuropharmacology, University of California, Los Angeles, CA, USA

**V. Ortiz** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

**Catarina R. Oliveira** Center for Neuroscience and Cell Biology (CNC) and Faculty of Medicine, University of Coimbra, Coimbra, Portugal

**María Carmen Ovejero-Benito** Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC), Madrid, Spain

**Björn Owe-Larsson** Department of Clinical Neuroscience, Karolinska Institutet, Section of Psychiatry at Karolinska University Hospital Huddinge, Stockholm, Sweden

Sophia T. Papadeas Chapel Hill, NC, USA

**Irmgard Paris** Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

Department of Basic Sciences, Santo Tomas University, Viña del Mar, Chile

Livia Pasquali Clinical Neurology, University of Pisa, Pisa, Italy

**Justyna Paterak** Department of Pharmacology and Physiology of the Nervous System, Institute of Psychiatry and Neurology, Warsaw, Poland

**R. Perez** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

Anabel Pérez-Gómez Department of Biomedical Sciences, University of Prince Edward Island, Charlottetown, PEI, Canada

**Marzia Perluigi** Department of Biochemical Sciences, Faculty of Pharmacy and Medicine, Sapienza University of Rome, Rome, Italy

**George Perry** Department of Biology and UTSA Neurosciences Institute, The University of Texas at San Antonio, San Antonio, TX, USA

Adam Perzyński Department of Psychiatry, Medical University of Lublin, Lublin, Poland

Laura Petrosini Department of Psychology, Sapienza University of Rome, Rome, Italy

I.R.C.C.S. Santa Lucia Foundation, Rome, Italy

Andrzej Pilc Institute of Pharmacology, Polish Academy of Sciences and Center of Excellence in Neuropsychopharmacology, Kraków, Poland

Jagiellonian University Medical College, Kraków, Poland

Juan C. Piña-Crespo Del E. Webb Center for Neuroscience, Aging, and Stem Cell Research, Sanford-Burnham Medical Research Institute, La Jolla, CA, USA

Annalisa Pinna Department of Biomedical Sciences, University of Cagliari, Cagliari, Italy

CNR Institute of Neuroscience, Cagliari, Italy

Augusta Pisanu CNR Institute of Neuroscience, Cagliari, Italy

Elisa Pizarro-Estrella Departamento de Bioquímica y Biología Molecular y Genética, Facultad de Enfermería y TO, Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas (CIBERNED), Universidad de Extremadura, Cáceres, Spain

Krzysztof Polberg Department of Otolaryngology, MSW Hospital, Lublin, Poland

Anne Poljak School of Medical Sciences, Faculty of Medicine and Bioanalytical Mass Spectrometry Facility, University of New South Wales, Sydney, Australia

**Bartosz Pomierny** Department of Toxicology, Jagiellonian University Medical College, Kraków, Poland

Lucyna Pomierny-Chamioło Department of Toxicology, Faculty of Pharmacy, Jagiellonian University, Kraków, Poland

**Dean L. Pountney** School of Medical Science and Griffith Health Institute, Griffith University, Gold Coast, QLD, Australia

**David V. Pow** School of Medical Sciences and Health Innovations Research Institute, RMIT University, Melbourne, VIC, Australia

**Pierre-François Pradat** Département des Maladies du Système Nerveux, AP-HP, Groupe Hospitalier Pitié-Salpêtrière, UMR-678, INSERM-UPMC, Paris, France

**Rui Daniel Prediger** Laboratório Experimental de Doenças Neurodegenerativas, Departamento de Farmacologia, Universidade Federal de Santa Catarina (UFSC), Florianópolis, SC, Brazil Paolo Preziosi Institute of Pharmacology, Catholic University School of Medicine, Rome, Italy

Robson Luiz Puntel Universidade Federal do Pampa, Uruguaiana, RS, Brazil

Rowan Radford Griffith Health Institute, Griffith University, Gold Coast, QLD, Australia

Lotten Ragnarsson Institute for Molecular Bioscience, The University of Queensland, Brisbane, QLD, Australia

Abdur Rahman Department of Food Science and Nutrition, College for Women, Kuwait University, Safat, Kuwait

**Rita Raisman-Vozari** INSERM UMR 975 (ex U679), CNRS UMR 7225, Université Pierre et Marie Curie (UPMC), Centre de Recherche de l'Institut du Cerveau et de la Moelle épinière (CRICM), Thérapeutique Expérimentale de la neurodégénérescence, Hôpital de la Salpêtrière, Paris, France

Matt S. Ramer International Collaboration on Repair Discoveries, The University of British Columbia, Vancouver, BC, Canada

Max Rapp-Ricciardi Department of Psychology, University of Gothenburg, Gothenburg, Sweden

**A. Cristina Rego** Center for Neuroscience and Cell Biology (CNC) and Faculty of Medicine, University of Coimbra, Coimbra, Portugal

**Serafino Ricci** Department of Anatomy, Histology, Forensic Medicine and Orthopaedics, Sapienza University of Rome, Rome, Italy

**Peter Riederer** University of Würzburg, Department of Psychiatry, Psychosomatics and Psychotherapy, and National Parkinson Foundation Centers of Excellence for Neurodegenerative Diseases Research, Laboratory for Clinical Neurochemistry, Würzburg, Germany

**B. Rivera** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

João Batista Teixeira Rocha Universidade Federal de Santa Maria, RS, Brazil

**Claudia Rodríguez-Ruiz** Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC), Madrid, Spain

**Mary-Louise Rogers** Department of Human Physiology, Centre for Neuroscience, School of Medicine, Flinders University, Adelaide, SA, Australia

**E. Rojas-Mancilla** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

Radosław Rola Department of Neurological Surgery, Medical University of Lublin, Lublin, Poland

Department of Physiopathology, Institute of Agricultural Medicine, Lublin, Poland

Marina Romero-Ramos Department of Biomedicine, Aarhus University, Aarhus C, Denmark

**C. M. Ros** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain

**F. Ros-Bernal** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain

**Elyse Rosa** Department of Psychiatry and Behavioural Neurosciences, McMaster University, Hamilton, ON, Canada

**Bartłomiej Rospond** Department of Toxicology, Faculty of Pharmacy, Jagiellonian University, Kraków, Poland

Khalil Saadipour Division of Health Sciences, University of South Australia, Adelaide, SA, Australia

**Perminder Sachdev** Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

Neuropsychiatric Institute, Prince of Wales Hospital, Sydney, Australia

**Anna Sadakierska-Chudy** Laboratory of Drug Addiction Pharmacology, Institute of Pharmacology Polish Academy of Sciences, Kraków, Poland

Sara Saez-Atienzar Faculty of Medicine and Dentistry, Catholic University of Valencia San Vicente Mártir, Valencia, Spain

**T. Sai** Department of Veterinary Pathology, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan

**M. Salkovic-Petrisic** Department of Pharmacology and Croatian Institute for Brain Research, School of Medicine, University of Zagreb, Zagreb, Croatia

**Thomas E. Salt** Department of Visual Neuroscience, UCL Institute of Ophthalmology, London, UK

**A. Sampedro** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

**Sara Sanz-Blasco** Del E. Webb Center for Neuroscience, Aging, and Stem Cell Research, Sanford-Burnham Medical Research Institute, La Jolla, CA, USA

**Arne Schousboe** Department of Drug Design and Pharmacology, Faculty of Health and Medical Sciences, University of Copenhagen, Copenhagen, Denmark

Angela L. M. Scott International Collaboration on Repair Discoveries, The University of British Columbia, Vancouver, BC, Canada

**Juan Segura-Aguilar** Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

**Y. Shibui** Department of Veterinary Pathology, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan

**Sohel Shivji** Department of Psychiatry and Behavioural Neurosciences, McMaster University, Hamilton, ON, Canada

Nicola Simola Department of Biomedical Sciences, University of Cagliari, Cagliari, Italy

**Bal Ram Singh** National Botulinum Research Center and Department of Chemistry and Biochemistry, University of Massachusetts Dartmouth, Dartmouth, MA, USA

Irena Smaga Department of Toxicology, Jagiellonian University Medical College, Kraków, Poland

**Maria E. Solesio** Unidad de Neuropsicofarmacología Traslacional, Complejo Hospitalario Universitario de Albacete, Albacete, Spain

Kim A. Staats Laboratory of Neurobiology and Leuven Research Institute of Neurodegenerative Diseases (LIND), KU Leuven, Leuven, Belgium

Vesalius Research Center, VIB, Leuven, Belgium

**Roman Stefański** Department of Pharmacology and Physiology of the Nervous System, Institute of Psychiatry and Neurology, Warsaw, Poland

Sarah Stegeman College of Medicine, University of Central Florida, Orlando, FL, USA

Andrzej Stepulak Department of Biochemistry and Molecular Biology, Medical University of Lublin, Lublin, Poland

Department of Otolaryngology, MSW Hospital, Lublin, Poland

**Trevor W. Stone** Institute of Neuroscience and Psychology, University of Glasgow, Glasgow, UK

**S. Subash** Department of Food Science and Nutrition, College of Agriculture and Marine Sciences, Sultan Qaboos University, Muscat, Oman

Kiminobu Sugaya College of Medicine, University of Central Florida, Orlando, FL, USA

**Ying Sun** School of Pharmacy and Medical Sciences, Faculty of Health Sciences, University of South Australia, Adelaide, SA, Australia

Levente Szalárdy Department of Neurology, University of Szeged, Szeged, Hungary

Arpad Szallasi Monmouth Medical Center, Long Branch, NJ, USA

**Henry Szechtman** Department of Psychiatry and Behavioural Neurosciences, McMaster University, Hamilton, ON, Canada

Fei Tan Department of Neurology, Shengjing Hospital of China Medical University, Shenyang, China

**E. Tarragón** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain

**R. Andrew Tasker** Department of Biomedical Sciences, University of Prince Edward Island, Charlottetown, PEI, Canada

**Carol J. Thiele** Cellular and Molecular Biology, Pediatric Oncology Branch, National Cancer Institute, National Institute of Health, Bethesda, MD, USA

József Toldi Department of Physiology, Anatomy and Neuroscience, University of Szeged, Szeged, Hungary

Neuroscience Research Group of the Hungarian Academy of Sciences and University of Szeged, Szeged, Hungary

Gabrielle N. Turski 2nd Faculty of Medicine, Charles University, Prague, Czech Republic

Michael Tymianski Neuroprotection Laboratory, Toronto Western Research Institute, University Health Network, Toronto, ON, Canada

**K. Uchida** Department of Veterinary Pathology, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan

**Ewa M. Urbańska** Laboratory of Cellular and Molecular Pharmacology, Department of Experimental and Clinical Pharmacology, Medical University of Lublin, Lublin, Poland

Department of Toxicology, Institute of Agricultural Medicine, Lublin, Poland

**Pamela Urrutia** Department of Biology, Faculty of Sciences, University of Chile, Casilla, Santiago, Chile

**J. L. Valdes** Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

**Ludo Van Den Bosch** Laboratory of Neurobiology and Leuven Research Institute of Neurodegenerative Diseases (LIND), KU Leuven, Leuven, Belgium

Vesalius Research Center, VIB, Leuven, Belgium

László Vécsei Department of Neurology, University of Szeged, Szeged, Hungary

Neuroscience Research Group of the Hungarian Academy of Sciences and University of Szeged, Szeged, Hungary

Sandra M. Vetiska Neuroprotection Laboratory, Toronto Western Research Institute, University Health Network, Toronto, ON, Canada

James C. Vickers Wicking Dementia Research and Education Centre, University of Tasmania, Hobart, Tasmania, Australia

**R. K. Vijayan** Department of Food Science and Nutrition, College of Agriculture and Marine Sciences, Sultan Qaboos University, Muscat, Oman

**Miquel Vila** Neurodegenerative Diseases Research Group, Vall d'Hebron Research Institute–CIBERNED, Barcelona, Spain

Department of Biochemistry and Molecular Biology, Autonomous University of Barcelona (UAB), Barcelona, Spain

Catalan Institution for Research and Advanced Studies (ICREA), Barcelona, Spain

Marta Volosin Department of Biological Sciences, Rutgers, The State University of New Jersey, Newark, NJ, USA

**Christopher S. von Bartheld** Department of Physiology and Cell Biology, University of Nevada School of Medicine, Reno, NV, USA

Steve Vucic Sydney Medical School Westmead, University of Sydney, Sydney, Australia

Neuroscience Research Australia, Randwick, Sydney, NSW, Australia

Matthew C. Walker UCL Institute of Neurology, University College London, London, UK

Yan-Jiang Wang Department of Neurology, Daping Hospital, Third Military Medical University, Daping, Chongqing, China

Mathew Wong Griffith Health Institute, Griffith University, Gold Coast, QLD, Australia

Erik Z. Woody Department of Psychology, University of Waterloo, Waterloo, ON, Canada

Junhua Xiao Department of Anatomy and Neuroscience, The University of Melbourne, Parkville, VIC, Australia

**Xingshun Xu** Department of Neurology, The Second Affiliated Hospital of Soochow University, Suzhou City, Jiangsu, China

The Institute of Neuroscience, Soochow University, Suzhou City, Jiangsu, China

**Manoj Yadav** National Institute of Science Education and Research, Institute of Physics Campus, Bhubaneswar, Orissa, India

**Hongyan Yang** David Geffen School of Medicine, Department of Chemistry and Biochemistry, Semel Institute for Neuroscience and Human Behavior, and Hatos Center for Neuropharmacology, University of California, Los Angeles, CA, USA

**Miao Yang** School of Pharmacy and Medical Sciences, Faculty of Health Sciences, University of South Australia, Adelaide, SA, Australia

**Reiji Yoshimura** Department of Psychiatry, School of Medicine, University of Occupational and Environmental Health, Yahatanishi-ku, Kitakyushu, Japan

**J. E. Yuste** Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain

Dénes Zádori Department of Neurology, University of Szeged, Szeged, Hungary

Fan Zeng Department of Neurology, Daping Hospital, Third Military Medical University, Daping, Chongqing, China

Xin-Fu Zhou School of Pharmacy and Medical Sciences, Faculty of Health Sciences, University of South Australia, Adelaide, SA, Australia

Part I

Neurotoxins

# **Survey of Selective Neurotoxins**

# Richard M. Kostrzewa

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R.M. Kostrzewa

Department of Biomedical Sciences, Quillen College of Medicine, East Tennessee State University, Johnson City, TN, USA e-mail: KOSTRZEW@mail.etsu.edu

## Abstract

There has been an awareness of nerve poisons from ancient times. At the dawn of the twentieth century, the actions and mechanisms of these poisons were uncovered by modern physiological and biochemical experimentation. However, the era of selective neurotoxins began with the pioneering studies of R. Levi-Montalcini through her studies of the neurotrophin "nerve growth factor" (NGF), a protein promoting growth and development of sensory and sympathetic noradrenergic nerves. An antibody to NGF, namely, anti-NGF – developed in the 1950s in a collaboration with S. Cohen – was shown to produce an "immunosympathectomy" and virtual lifelong sympathetic denervation. These Nobel Laureates thus developed and characterized the first identifiable selective neurotoxin. Other selective neurotoxins were soon discovered, and the compendium of selective neurotoxins, with the potential to destroy or produce dysfunction of a variety of phenotypic nerves.

Selective neurotoxins are of value because of their ability to selectively destroy or disable a common group of nerves possessing (1) a particular neural transporter, (2) a unique set of enzymes or vesicular transporter, (3) a specific type of receptor or (4) membranous protein, or (5) other uniqueness.

The era of selective neurotoxins has developed to such an extent that the very definition of a "selective" neurotoxin has warped. For example, (1) N-methyl-Daspartate receptor (NMDA-R) antagonists, considered to be neuroprotectants by virtue of their prevention of excitotoxicity from glutamate receptor agonists, actually lead to the demise of populations of neurons with NMDA receptors, when administered during ontogenetic development. The mere lack of natural excitation of this nerve population, consequent to NMDA-R block, sends a message that these nerves are redundant - and an apoptotic cascade is set in motion to eliminate these nerves. (2) The rodenticide rotenone, a global cytotoxin that acts mainly to inhibit complex I in the respiratory transport chain, is now used in low dose over a period of weeks to months to produce relatively selective destruction of substantia nigra dopaminergic nerves and promote alpha-synuclein deposition in brain to thus model Parkinson's disease. Similarly, (3) glial toxins, affecting oligodendrocytes or other satellite cells, can lead to the damage or dysfunction of identifiable groups of neurons. Consequently, these toxins might also be considered as "selective neurotoxins," despite the fact that the targeted cell is nonneuronal. Likewise, (4) the dopamine D<sub>2</sub>-receptor agonist quinpirole, administered daily for a week or more, leads to development of D2-receptor supersensitivity - exaggerated responses to the D<sub>2</sub>-receptor agonist, an effect persisting lifelong. Thus, neuroprotectants can become "selective" neurotoxins; nonspecific cytotoxins can become classified as "selective" neurotoxins; and receptor agonists, under defined dosing conditions, can supersensitize and thus be classified as "selective" neurotoxins. More examples will be uncovered as the area of selective neurotoxins expands.

The description and characterization of selective neurotoxins, with unmasking of their mechanisms of action, have led to a level of understanding of neuronal activity and reactivity that could not be understood by conventional physiological observations. This chapter will be useful as an introduction to the scope of the field of selective neurotoxins and provide insight for in-depth analysis in later chapters with full descriptions of selective neurotoxins.

# Keywords

3-Nitropropionic acid • 5,6-Dihydroxytryptamine • 5,7-Dihydroxytryptamine • 6-Hydroxydopa • 6-Hydroxydopamine • AF64A • Amphetamine • AMPA • AOAA • BMAA • BOAA • Botulinum neurotoxin • Capsaicin • Cocaine • Domoic acid • DSP-4 • Excitatory amino acids • Glutamic acid • Haloperidol • IgG-Saporin • Kainic acid • Kynurenine • Methamphetamine • Methylenedioxymethamphetamine • MPTP • Neurotoxins • Parachloroamphetamine • Quinolinic acid • Quinpirole • Vanilloids

List of Abbreviations				
3-NP	3-Nitropropionic acid			
5,6-DHT	5,6-Dihydroxytryptamine			
5,7-DHT	5,7-Dihydroxytryptamine			
6-OHDA	6-Hydroxydopamine			
6-OHDOPA	6-Hydroxydopa			
AF64A	Ethylcholine aziridinium ion			
AMPA	α-amino-2,3-hydroxy-5-methyl-3-oxo-isoxazole-propionic acid			
AMPH	Amphetamine			
AOAA	aminooxyacetic acid			
BMAA	β-N-methylamino-L-alanine			
BOAA	β-N-oxalylamino-L-alanine			
BoNT	Botulinum neurotoxin			
DSP-4	2-chloroethylamines N-(2-chloroethyl)-N-ethyl-2-bromobenzylamine			
EAAs	excitatory amino acids			
GLU	Glutamic acid			
HPTP	4-(4-chlorophenyl)-1,4-(4-fluorophenyl)-4-oxobutyl-1,2,3,6-			
	tetahydropyridine			
MDMA	Methylenedioxymethamphetamine			
METH	Methamphetamine			
MPTP	1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine			
PCA	Parachloroamphetamine			
$RHPP^+$	4-(4-chlorophenyl)-1,4-(4-fluorophenyl)-4-hydroxybutyl-			
	pyridinium (see)			

# 1 Introduction

The renowned physiologist Walter B. Canon promoted the principle that the function of a nerve can be deduced by denoting the deficits produced when that nerve is destroyed. The implication is that the nerve normally acts in a way to promote functions that prevent the "absence deficits." This reasoning is sound, and this experimental approach has been proven to be invaluable. However, while it is practical to section axons and while certain neuronal nuclei can be electrolytically destroyed with only minor injury to surrounding nerves, this approach is not so useful for brain and spinal cord because of the unintended and consequential destruction of neuronal nuclei and axons, particularly if attempting to surgically disrupt deep brain nuclei.

The advantage of selective neurotoxins is that these agents can target specific neuronal phenotypes while leaving surrounding tissue unaffected and while leaving virtually all other nerve types intact. It matters not that the target group of neurons is deep in the brain or dispersed in the spinal cord. In this way, the "absence deficit" can be discerned, and the normal function of this population of nerves can be deduced. This experimental approach is now commonplace.

Another arena for selective neurotoxins is in the modeling of neurological and psychiatric disorders. The neurotoxin 6-hydroxydopamine (6-OHDA), possessing high affinity for the norepinephrine (NE) transporter (NET), has been used to produce sympathectomy in experimental animals (Thoenen and Tranzer 1968) and also used to model attention deficit hyperactivity disorder (ADHD) (Shaywitz et al. 1976) and Lesch-Nyhan syndrome (Breese et al. 1984b). The neurotoxin 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) has been used to model Parkinson's disease (Langston and Ballard 1984). N-methyl-D-aspartate receptor (NMDA-R) antagonists have been used to model schizophrenia (Farber et al. 1995; see Coyle 1996). More recently, repeated treatments with the dopamine (DA) D<sub>2</sub>-receptor agonist quinpirole have likewise modeled schizophrenia (Peng et al. 1990; see Brown et al. 2012) as well as obsessive-compulsive disorder (Szechtman et al. 1998). Reversible and permanent models of tardive dyskinesia have been produced by haloperidol in intact and in 6-OHDA-lesioned rats, respectively (Waddington et al. 1983; Huang et al. 1997b). Quinolinic acid lesions have modeled Huntington's disease (Schwarcz et al. 1984; Beal et al. 1988). Neurotoxins have been used to model additional neurological and psychiatric disorders, and the list will continue to grow with time.

With anti-NGF having been developed in the 1950s and with both 6-hydroxydopamine and glutamate having been used as neurotoxins since the 1960s, the era of selective neurotoxins is 6 decades in the making. During the twenty-first century, the number of selective neurotoxins will inevitably grow, thereby expanding the potential and usefulness of these agents in increasing our understanding of the processes and mechanisms entailed in neuronal function. Additionally, selective neurotoxins will model ever more neurological and psychiatric disorders and thereby improve the template for development of better treatments for human disorders.

# 2 Terminology

The term "selective neurotoxin" refers to an agent which has toxic potential and relative selectivity for a neuron with:

- 1. A defined phenotype (e.g., dopaminergic nerve)
- 2. A specified receptor population which confers susceptibility to damage by agonists (e.g., glutamate-induced excitotoxicity)

- 3. A cellular protein that is inactivated by a particular agent (e.g., DSP-4 and noradrenergic nerves)
- 4. A metabolic schema that is specifically inactivated by an agent (e.g., rotenone inhibition of mitochondrial complex I, leading to degeneration of dopaminergic nerves)
- 5. A protein, uniquely targeted by an immunotoxin (e.g., IgG-saporin and cholinergic nerves)
- 6. Surface protein targeted by an agent that renders the nerve inactive (e.g., botulinum neurotoxin and cholinergic nerves)
- 7. Ion channels rendered inactive by an agent (e.g., tetrodotoxin and sodium channels)

Agents and antibodies directed towards neurotrophins can similarly prevent development of a population of neurons which are dependent on neurotrophic actions particularly at defined stages in ontogeny (e.g., anti-nerve growth factor [anti-NGF] and paravertebral sympathetic noradrenergic nerves).

As more elements in nerves become targets, the list above will expand with time. And with time, the definition of a neurotoxin will blur. Already recognized is the fact that NMDA-R antagonists can result in a neurotoxic effect by virtue of their neuroprotective action (Olney et al. 1991; Farber et al. 1995). Also, when administered particularly in ontogeny, NMDA-R antagonists prevent neural excitation by endogenous glutamate – thereby sending a "message" that the now quiescent population of neurons is unnecessary. This group of developing neurons dies, and the net effect of the neuroprotective antagonist is overt neurotoxicity (see Olney et al. 2002).

In another vein, repeated treatments with some agonists – notably the DA  $D_2$ -receptor agonist quinpirole – result in prolonged or even permanent supersensitization of  $D_2$  receptors. Following the development of receptor supersensitivity, animals when challenged acutely with quinpirole display exaggerated behaviors; and this disposition to a  $D_2$  agonist is lifelong (Kostrzewa and Brus; 1991; Kostrzewa 1995; Kostrzewa et al. 2003). Thus, in the absence of overt neuronal cell loss, and with no discernible anatomical effect, altered behavioral outcome represents an abnormal response and thereby an index of neurotoxicity.

A neurotoxin thus can be classified as such, even in the absence of neuronal apoptosis or necrosis. Therefore, a selective neurotoxin can be defined only loosely, and the definition surely will be more ambiguous over time. Nevertheless, the utility and advantages of neurotoxins will grow over the next century and beyond, and those who employ selective neurotoxins will recognize this spectrum of agents by outcomes in the experimental realm.

The topic of selective neurotoxins has been reviewed recently (Kostrzewa 1999b, 2009; Segura-Aguilar and Kostrzewa 2004; 2006; Herrera-Marschitz et al. 2007).

# 3 Early History of Selective Neurotoxins

The stage for selective neurotoxins was constructed in the 1930s by the noted physiologist Walter B. Canon, who promoted the thesis that the function of

a nerve can be deduced by denoting altered effect when the nerve is destroyed by surgery or other physical means (Cannon and Rosenblueth 1937). While this thesis has proven to be valuable, limitations of a surgical experimental approach relate to the fact that multiple types of neurons [phenotypically different (i.e., having different neurotransmitter systems) or with different populations of receptor types] are simultaneously damaged. Accordingly, it is not possible to ascribe a physiological alteration to a phenotypically distinct type of neuron.

The initial discovery of a selective neurotoxin was conceivably that by Lucas and Newhouse (1957) who found that the amino acid glutamate (Glu) produced overt destruction of neurons in immature mouse retina. Years later this effect was attributed to excess neuronal excitation, resulting in "excitotoxicity."

However, the age of selective neurotoxins arrived with the seminal work of the Nobel Laureate Rita Levi-Montalcini. During her fellowship with Viktor Hamburger in the 1950s, the discovery was made that there is a factor in mouse sarcoma 180 and 37, also in snake venom and mouse salivary glands, that profoundly promotes the growth of sensory and sympathetic ganglia. In collaboration with the biochemist Stanley Cohen, this neurotrophin was ultimately isolated and termed "nerve growth factor" (NGF) (Cohen et al. 1956; Levi-Montalcini et al. 1954; Levi-Montalcini and Cohen 1956; Cohen and Levi-Montalcini 1957; Levi-Montalcini and Booker 1960b; Levi-Montalcini 1964). Subsequently, an antiserum to NGF was developed and shown – when administered to neonatal mice – to prevent the development of sympathetic (noradrenergic) ganglia, thereby resulting in a relative noradrenergic denervation of peripheral organs. While intermediary ganglia (e.g., mesenteric ganglion) and terminal ganglia remained partially or largely intact, paravertebral ganglia were near-totally destroyed. The outcome was predominate removal of the sympathetic nervous system - an "immunosympathectomy" (Levi-Montalcini and Booker 1960a; Levi-Montalcini 1966; 1987 (reviews); Levi-Montalcini and Angeletti 1966), R. Levi-Montalcini and S. Cohen shared the 1987 Nobel Prize for their work. Anti-NGF ushered in the era of selective neurotoxins.

While anti-NGF was a milestone in providing discrete denervation of a phenotypically selective class of nerves, anti-NGF was not a neurotoxin per se. Rather, anti-NGF prevented the action of a specific neurotrophin, which thus prevented programmed ontogenetic development of sympathetic noradrenergic nerves. Anti-NGF had limitations: (1) Noradrenergic fibers originating in intermediary sympathetic ganglia (e.g., mesenteric ganglia) partially survived the effects of anti-NGF, and (2) sympathetic terminal ganglia, largely to genitourinary regions, largely survived anti-NGF treatment. As an experimental strategy, anti-NGF (3) had virtually no discernible effect in adult species, and (4) it had no effect on noradrenergic nerves in the brain. Anti-NGF thus was the prologue of agents that are specifically classified as selective neurotoxins.

# 4 The Era of Synthetic and Small Molecule Selective Neurotoxins

Beginning in the 1960s and into the present, there has been a series of discoveries of small molecule selective neurotoxins. These agents represent a valuable resource towards determining the roles of identifiable (i.e., sometimes phenotypically distinct) neurons in physiological and behavioral processes. Moreover, these neurotoxins have been useful likewise in producing animal models of human psychiatric, neurologic, and neurodegenerative disorders. The brief survey of selective neurotoxins herein provides a brief history of discovery as well as a description of their effects and resulting utility.

# 5 Survey of Individual Selective Neurotoxins

# 5.1 6-Hydroxydopamine (6-OHDA)

## 5.1.1 Historical Perspective

Porter et al. (1963; 1965) and Stone et al. (1963; 1964) in their mutually collaborative studies made the discovery that the compound 2,4,5-trihydroxyphenethylamine (6-hydroxydopamine, 6-OHDA) produced long-lived depletion of NE in the heart and other tissues. However, its mechanism of action was undefined. Several years later, in search of an electrophilic marker for electron microscopic visualization of synaptic granules in sympathetic neurons, Hans Thoenen and J.P. Tranzer found a near ideal candidate in 5-hydroxydopamine (5-OHDA) (Tranzer and Thoenen 1967a). However, the startling finding was that the analog 6-OHDA produced overt neurotoxicity, destroying sympathetic (noradrenergic) neurons (Tranzer and Thoenen 1967b; 1968; Thoenen and Tranzer 1968). In short time, it was shown that 6-OHDA, when injected into brain tissue or into a brain ventricle, destroyed dopaminergic as well as noradrenergic neurons (Bloom et al. 1969; Uretsky and Iversen 1969; 1970). Many facets of 6-OHDA actions were described in review articles, shortly after discovery of 6-OHDA neurotoxicity (Thoenen and Tranzer 1973; Kostrzewa and Jacobowitz 1974). The key terms "6-hydroxydopamine" or "6-OHDA" appear in 10,846 papers in PubMed, as of January 30, 2013. 6-OHDA is still widely used today.

## 5.1.2 Selectivity of Action

6-OHDA is a relatively selective neurotoxin for noradrenergic and dopaminergic neurons. Owing to its chemical similarity to DA (Fig. 1), 6-OHDA (Fig. 2) has high affinity for the NET and DA transporter (DAT) and entry into NE neurons and DA neurons resulting in a high intraneuronal concentration of 6-OHDA. The K<sub>i</sub> of 6-OHDA for the NET is  $<50 \mu$ M (Iversen 1970; Ma et al. 1995). When selective damage to dopaminergic nerves is the objective, an NET inhibitor such as desipramine is administered as pretreatment (Breese and Traylor 1971; Peters et al. 1977). During the process of uptake by the NET, 6-OHDA releases NE to produce a sympathomimetic effect (De Champlain and Nadeau 1971; De Champlain and Van Amerigen 1972).



**Fig. 2** (From Kostrzewa 1999b)

# 5.1.3 Mechanism of Action

Because 6-OHDA has micromolar affinity for the vesicular monoamine transporter-2 (VMAT-2) on NE granules and DA granules (Jonsson and Sachs 1975; Sachs et al. 1975), 6-OHDA is accumulated by granules and stored as a false transmitter.

Inside the nerve, 6-OHDA auto-oxidizes to a number of intraconvertible reactive oxygen species (ROS) including 6-OHDA-quinones, aminochrome, 5,6dihydroxyindole, indolines, and indole-quinones (Senoh and Witkop 1959; Saner and Thoenen 1971a, b; see Kostrzewa and Jacobowitz 1974). The destructive role of 6-OHDA-derived quinoid species is likely due not so much to their binding to cellular elements, but rather to their generation of a family of other ROS which overwhelms the endogenous cellular neuroprotective processes (Graham et al. 1978). 6-OHDA-ortho-quinone and 6-OHDA-para-quinone are resonant tautomers, each of which is capable of destroying noradrenergic nerves (Heikkila et al. 1973a). However, it appears that 6-OHDA-o-semiquinone is the paramount neurotoxic auto-oxidative species producing neurotoxicity (Segura-Aguilar 2001).

Most of the above ROS generate intraneuronal hydrogen peroxide  $(H_2O_2)$ , which itself is toxic (Heikkila and Cohen 1971, 1972a, b). Intraneuronal catalase inactivates  $H_2O_2$ , converting it to water. Superoxide radical also is generated by 6-OHDA auto-oxidation (Heikkila and Cohen 1973; Cohen and Heikkila 1974), as well as highly reactive hydroxyl radical (Cohen and Heikkila 1974).

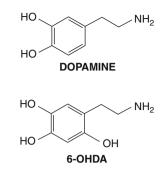
# 5.1.4 Effects of 6-OHDA Treatment of Adult Rodents

Being ionized, 6-OHDA is unable to cross the blood-brain barrier (BBB). If administered peripherally, 6-OHDA produces marked sympathetic denervation to most peripheral organs.

To produce a CNS effect in adults, 6-OHDA must be administered into a cerebral spinal fluid compartment (i.e., intraventricular, intracisternal, or intrathecal spaces) or directly into CNS parenchyma (e.g., intrastriatal, intranigral).

# 5.1.5 Intraparenchymal 6-OHDA Treatments in Rat Brain

Direct injection of 6-OHDA into substantia nigra pars compacta (SNpc) – to model Parkinson's disease (PD) – results in profound (>90 %) destruction of dopaminergic perikarya, with associated near-total dopaminergic denervation of neostriatum (Ungerstedt 1968, 1971a). Because such bilateral 6-OHDA lesions are accompanied



by aphagia, adipsia, immobility, lack of grooming, and ultimate death, intranigral 6-OHDA lesions are commonly given unilaterally. Rats with a unilateral intranigral 6-OHDA lesion are able to eat, drink, groom, and otherwise remain mobile but with a circling towards the lesioned side, which abates to a large degree over time (Ungerstedt 1971d).

Intrastriatal 6-OHDA administration results in an acute moderate destruction of dopaminergic striatal terminals, followed by progressive dopaminergic denervation of this brain region over a period of 4 months, with ultimate death of 50–70 % of SNpc dopaminergic perikarya (Berger et al. 1991; Sauer and Oertel 1994; Przedborski et al. 1995; Blandini et al. 2007).

6-OHDA directly administered into the locus coeruleus results in noradrenergic denervation of neocortex, hippocampus, cerebellum, and other regions, to a lesser extent (Ungerstedt 1971c).

# 5.1.6 Intracisternal (i.c.) and Intraventricular (i.vtr.) 6-OHDA Treatments of Rats

In adult rats i.c. and i.vtr. 6-OHDA treatments produce modest effects on noradrenergic and dopaminergic innervation of the brain, with potency increased by pretreatment with a monoamine oxidase inhibitor (MAO-I) and with selectivity for dopaminergic neurons increased by pretreatment with the NET inhibitor desipramine (see *Handbook of Neurotoxicity* chapter by Papadeas and Breese 2013). DA D<sub>2</sub>receptor (D<sub>2</sub>-R) supersensitivity is prominent in rats so-lesioned (Ungerstedt 1971b).

## 5.1.7 Effects of 6-OHDA Treatment of Neonatal Rodents

When administered non-centrally to perinatal rodent species, 6-OHDA and related neurotoxins have a dual action in destroying peripheral sympathetic noradrenergic nerves. 6-OHDA initially destroys sympathetic noradrenergic nerves via generation of intraneuronal ROS, and this effect results in the inability of noradrenergic nerve endings to enact the effects of NGF which would promote further ontogenetic development of these nerves. In perinatal rats, 6-OHDA crosses the blood-brain barrier and produces noradrenergic neuronal degeneration in the brain. When administered to rats on the day of birth, 6-OHDA (60 mg/kg ip) produces lifelong partial noradrenergic denervation of spinal cord (~75 % decrease) and neocortex  $(\sim 50\%)$ , with eventual noradrenergic sprouting and hyperinnervation of midbrain ( $\sim$ 50 %), pons-medulla ( $\sim$ 50 % increase), and cerebellum ( $\sim$ 100 % increase). When administered up to 5 days post-birth, 6-OHDA results in an approximate 40 % permanent decrease in noradrenergic innervation of hippocampus; when given from 3 days to 14 days post-birth, 6-OHDA decreases noradrenergic innervation of cerebellum by  $\sim$ 50 %. The blood-brain barrier forms in different parts of the brain at different postnatal times, accounting for 6-OHDA access to different brain regions at different postnatal times (Kostrzewa 2007).

Both i.c. and i.vtr. 6-OHDA treatments in perinatal rodents result in near-total destruction of SNpc (Berger et al. 1985; Fernandes Xavier et al. 1994) and associated near-total DA depletion (Breese et al. 1984a, b, 1985a, b, 1987; Bruno et al. 1987; Jackson et al. 1988; Dewar et al. 1990), with near-total dopaminergic

denervation of neostriatum (Snyder et al. 1986; Descarries et al. 1992). Selectivity of 6-OHDA for dopaminergic neurons is provided by desipramine pretreatment. In the absence of desipramine perinatal 6-OHDA largely destroys the noradrenergic system as well, with locus coeruleus perikarya being susceptible to damage. Notably, neonatal 6-OHDA lesioning of dopaminergic systems results in reactive serotoninergic sprouting and eventual 5-HT neuronal hyperinnervation of forebrain (Breese et al. 1984a, b; Stachowiak et al. 1984; Snyder et al. 1986; Luthman et al. 1987; Towle et al. 1989; Descarries et al. 1992).

In adult rats that are largely DA denervated by neonatal 6-OHDA treatment, the first challenge dose of a DA D<sub>1</sub>-R agonist induces marked perioral activity (vacuous chewing movements, VCMs), with the effect being dose related and with sensitivity to the DA D<sub>1</sub> agonist being 1,000-fold greater than observed in adult controls (Kostrzewa and Gong 1991). This effect is mediated or moderated by the serotoninergic system, as indicated by the fact that a 5-HT<sub>2</sub>-R antagonist attenuates the D<sub>1</sub> agonist effect (Gong et al. 1992) as does 5-HT fiber denervation produced by 5, 7-dihydroxytryptamine (5,7-DHT) treatment (Brus et al. 1994). This DA D<sub>1</sub>-R supersensitivity is accompanied by an increase in DA D<sub>1</sub>-receptor mRNA levels in neostriatum but without a change in the number of high-affinity D<sub>1</sub>-R in striatum (Gong et al. 1994; see Kostrzewa et al. 1998).

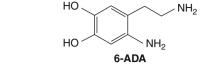
In the adult rats that were lesioned neonatally with 6-OHDA, the serotoninergic hyperinnervation of forebrain is accompanied by 5-HT-R supersensitivity, as evidenced by enhanced perioral activity in response to low-dose 5-HT<sub>2</sub> agonist treatment (Gong and Kostrzewa 1992; Gong et al. 1993).

For locomotor and classical stereotypic behaviors, there is latent  $D_1$ -R supersensitivity, as exemplified by the findings of Breese et al. (1985a, b, 1987), demonstrating that the first and second weekly adulthood challenge doses of a  $D_1$  agonist produce no enhanced behavioral response in the neonatally 6-OHDA-lesioned rats. However, the third and all subsequent doses of a  $D_1$ -R agonist produce profound locomotor and stereotypical responses – and the effect is lifelong (Breese et al. 1985a, b, 1987). The neonatally 6-OHDA-lesioned rat represents a unique animal model for studying DA-R supersensitivity (see Kostrzewa 1995; Papadeas and Breese (2013 – this Handbook of Neurotoxicity)).

The effects of 6-OHDA treatments of both adult rats and neonatal rats are discussed in several reviews (Kostrzewa and Jacobowitz 1974; Breese 1975; Zigmond and Stricker 1989; Breese et al. 1990, 1994, 2005; Zigmond et al. 1993; Breese and Breese 1998; Zigmond and Keefe 1998).

# 5.2 6-OHDA Analogs

A number of analogs of 6-OHDA [6-OHDA = 2,4,5-trihydroxyphenethylamine] have been described (Tranzer and Thoenen 1973), having effects similar to that of 6-OHDA: 6-aminodopamine (6-ADA) (Fig. 3) (Blank et al. 1972; Heikkila et al. 1973b; Jonsson and Sachs 1973); 2-methyl-6-OHDA (Lundstrom et al. 1973; Borchardt et al. 1977); 5-methyl-6-OHDA; 2,5-dimethyl-6-OHDA **Fig. 3** (From Kostrzewa 1999b)



(Borchardt et al. 1977); 4-amino-6-OHDA (Cheng and Castagnoli 1984); 2,3,5trihydroxyphenethylamine; 2,3,4,5-tetrahydrophenethylamine (Lundstrom et al. 1973); all eight of the trisubstituted phenyl analogs of phenethylamine; alphamethylphenethylamine; and 1-(5-amino-2,4-dihydroxyphenyl)-2-aminopropane (Ma et al. 1995). 6-OHDA analogs that form para-quinoidal species represent the neurotoxic forms (Cheng and Castagnoli 1984); however, the ease of oxidation is not directly correlated with neurotoxic potential (Ma et al. 1995). In early studies, 2,3,4-trihydroxy-; 2,3,6-trihydroxy-; 3,4,5-trihydroxy-; 2,4,6-trihydroxy-; and N, N-dimethyl-2,4,5-trihydroxyphenethylamine had no long-term effect on noradrenergic systems (Lundstrom et al. 1973).

# 5.3 6-Hydroxydopa (6-OHDOPA)

On the basis of knowledge that L-3,4-dihydroxyphenylalanine (L-DOPA) (Fig. 4) is effective as an anti-parkinsonian agent by virtue of its being able to cross the bloodbrain barrier, to then be converted in dopaminergic nerves to DA, 6-hydroxydopa (6-OHDOPA) (Fig. 5) was developed as a protoxin specifically for its ability to likewise cross the blood-brain barrier and then be converted to 6-OHDA and thereby destroy noradrenergic and/or dopaminergic nerves in the brain.

## 5.3.1 Historical Background

Ong et al. (1969) and Berkowitz et al. (1970) demonstrated that 6-OHDOPA effectively reduced NE content of the rat brain and that the detrimental effects of 6-OHDOPA on sympathetic noradrenergic nerves could be prevented by pretreatment with a dopa decarboxylase inhibitor. On the basis of these findings, RM Kostrzewa, under auspices of his dissertation advisor David M. Jacobowitz, undertook a series of biochemical and histochemical studies to determine the neurotoxic potential of 6-OHDOPA. Using a histofluorescence method to visualize noradrenergic neurons, the first evidence of 6-OHDOPA neurotoxicity was indicated by the loss of noradrenergic fiber number in the mouse brain, particularly forebrain – accompanied by the associated reduction in NE content in various brain regions (Jacobowitz and Kostrzewa 1971). Additionally, swollen preterminal axons were prominent in 6-OHDOPA-lesioned mice, thereby highlighting noradrenergic tracts – which ordinarily are not visible – and thereby facilitating the mapping of noradrenergic fiber tracts in the brain (Jacobowitz and Kostrzewa 1971).

Shortly thereafter, Sachs and Jonsson (1972a, b) reported analogous neurochemical findings, demonstrating NE reductions in the brains of 6-OHDOPA treated animals. While the advantage of 6-OHDOPA relates to its ability to be administered



non-centrally, 6-OHDOPA does actually produce neurotoxic effects when administered centrally (Richardson and Jacobowitz 1973). Definitive proof of 6-OHDOPA overt neurotoxicity was subsequently demonstrated by a decrease in the  $V_{max}$  for the NE transporter in brain (Kostrzewa and Garey 1976), loss of tyrosine hydroxylase activity in locus coeruleus (Kostrzewa et al. 1978), silver staining, and electron microscopic evidence of necrotic noradrenergic locus coeruleus neurons in the brain of 6-OHDOPA treated rats (Kostrzewa and Harper 1974, 1975). Cell counts of paraffin-embedded serial sections through the locus coeruleus established that neurons in the caudal half of the locus coeruleus (50 % cell loss) were most susceptible to 6-OHDOPA neurotoxicity (Clark et al. 1979).

# 5.3.2 Advantages of 6-OHDOPA Versus 6-OHDA

- 1. The protoxin 6-OHDOPA, unlike 6-OHDA, can be administered peripherally (ip, sc, iv) to adults to destroy noradrenergic nerves in the brain, thus avoiding direct intracisternal, intraventricular, or intraparenchymal injections of the brain.
- 2. Peripheral noradrenergic/sympathetic nerves can be protected and left virtually intact after 6-OHDOPA treatment by administering a peripherally acting dopa decarboxylase inhibitor, as pretreatment analogous to the conventional combination of a peripheral decarboxylase inhibitor with L-DOPA therapy of Parkinson's disease.
- 3. 6-OHDOPA is relatively selective for noradrenergic versus dopaminergic nerves.
- 4. Destruction of noradrenergic terminals results in retrograde accumulation of NE in swollen axons, which become clearly visible by a histofluorescence method for NE (see below), and thus facilitates the mapping of noradrenergic tracts in the brain.

# 5.3.3 Disadvantages of 6-OHDOPA Versus 6-OHDA

- 1. Because 6-OHDOPA is relatively selective for noradrenergic neurons, dopaminergic neurons are left largely intact and unaffected by 6-OHDOPA treatment.
- 2. 6-OHDOPA has low neurotoxic potency for noradrenergic neurons in adults. Consequently, it is not possible to produce near-total noradrenergic denervation of regions of the brain.

- 3. In high dose, 6-OHDOPA produces lethal effects, thus eliminating the possibility to use a large 6-OHDOPA dose.
- 4. 6-OHDOPA is much more expensive than 6-OHDA.

# 5.3.4 Effects in the Periphery and Brain of Rodents Lesioned as Adults with 6-OHDOPA

When administered peripherally to adults, 6-OHDOPA damages sympathetic noradrenergic terminals, thereby reducing sympathetic innervation of peripheral organs such as the heart and spleen, and this is accompanied by NE depletion of the same organs. In time, because sympathetic ganglia remain intact along with preterminal axons, nerve terminals to peripheral organs regenerate, and after a period of 4–8 weeks, most or all peripheral organs are fully reinnervated by sympathetic noradrenergic nerves (Kostrzewa and Jacobowitz 1972, 1974).

# 5.3.5 Long-Lived Effects in the Brain of Rodents Lesioned as Neonates with 6-OHDOPA

6-OHDOPA finds greatest utility when administered to *neonatal* rats or mice, in order to determine developmental effects on noradrenergic neurons and to study their regenerative capacity. In prenatal mice (Kostrzewa et al. 1978) and in postnatal rats, 6-OHDOPA produces long-lived near-total noradrenergic denervation of hippocampus and >85 % long-lived noradrenergic denervation of neocortex and also spinal cord (Kostrzewa 1975; Kostrzewa and Garey 1976; Kostrzewa and Jacobowitz 1973; Sachs and Jonsson 1972a, b; Zieher and Jaim-Etcheverry 1973, 1975). These regions are provided with noradrenergic innervation by axons originating from locus coeruleus, a nucleus prominently damaged by neonatal 6-OHDOPA treatment (Kostrzewa and Harper 1974; Clark et al. 1979). Noradrenergic innervation in ventral brain is less affected by 6-OHDOPA (e.g., hypothalamus and thalamus), and recovery in these regions occurs by 8 weeks (Kostrzewa and Garey 1976).

The pons-medulla and cerebellum of neonatally 6-OHDOPA-lesioned rats display noradrenergic hyperinnervation in adulthood, subsequent to reactive noradrenergic collateral fiber sprouting (Kostrzewa 1975; Kostrzewa and Garey 1976, 1977; Kostrzewa et al. 1978). Noradrenergic fiber sprouting in brain regions proximal to the locus coeruleus appears to be related to damage to the dorsal bundle and consequent noradrenergic denervation of forebrain (Jacobowitz and Kostrzewa 1971), because the effect is duplicated by non-pharmacological surgical transection of the dorsal bundle in neonatal rats – resulting in noradrenergic fiber hyperinnervation of regions proximal to the locus coeruleus, including the pons, medulla, and cerebellum (Klisans-Fuenmayor et al. 1986).

Sympathetic noradrenergic neurons are damaged by neonatal 6-OHDOPA, but recovery from damage – evidenced by return of NE content of the heart, spleen, and other organs – is complete by 8 weeks after birth. Dopaminergic innervation is virtually unaltered by neonatal 6-OHDOPA treatment, as evidenced by retention of normal DA content of striatum throughout the developmental period, up to 8 weeks and through 1 year and longer (Kostrzewa 1975; Kostrzewa and Garey 1976).

6-OHDOPA neurotoxicity is discussed in greater detail in several reviews (Kostrzewa and Jacobowitz 1974; Kostrzewa 1989, 1998).

# 5.4 DSP-4 and Xylamine

## 5.4.1 Overview

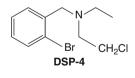
2-Chloroethylamine nitrogen mustards, N-2(2-chloroethyl)-N-ethyl-2-bromobenzylamine (DSP-4) (Fig. 6) and xylamine (Fig. 7) (Ross 1976; Ross and Reny 1976; Cho et al. 1980; Kammerer et al. 1979; Kreuger and Cook 1975) are able to cross the blood-brain barrier to selectively target noradrenergic neurons. These agents form quaternary aziridinium ions which have high affinity for the NET. Both covalently bind to the NET and are accumulated by noradrenergic neurons. These agents alkylate and inactivate proteins, thus effectively reducing the number of active NET sites (Fischer and Cho 1982) and impairing the function of intracellular cytochromes, enzymes (e.g., dopamine-beta-hydroxylase) (Ross 1976), and other proteins. In the process, NE stores are depleted in both the periphery and central nervous system (CNS) (Jonsson et al. 1981; Zieher and Jaim-Etcheverry 1980; Ross 1976) and overt noradrenergic neuronal destruction ensues (Fritschy and Grzanna 1989, 1991a, b; Fritschy et al. 1990; Grzanna et al. 1989; Jaim-Etcheverry and Zieher 1980; Zieher and Jaim-Etcheverry 1980).

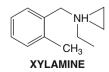
## 5.4.2 Mechanism of Action of DSP-4 and Xylamine

DSP-4 and xylamine are amines that intracyclize to form aziridinium ions (Ross et al. 1973) – nitrogen mustards (alkylating agents) that covalently and irreversibly bind to electrophilic sites such as thiols and amines in proteins, to effectively impair function of the protein and other molecules (Ranson et al. 1982). DSP-4 and xylamine have high affinity for the NET, and it is this property that confers selectivity of the molecules for noradrenergic neurons. Also, it is via the NET that DSP-4 and xylamine gain entry to the neurons. When DSP-4 binding to the NET is prevented by desipramine, the neurotoxic effect of DSP-4 is abated (Ross 1976). Although the precise mechanism is unknown, monoamine oxidase-B (MAO-B) inhibition by deprenyl (selegiline) also prevents DSP-4 neurotoxicity (Gibson 1987).

Fig. 6 (From Kostrzewa 1999b)

**Fig. 7** (From Kostrzewa 1999b)





However, because the MAO-B inhibitor MDL 72974 is not neuroprotective, it is not MAO-B inhibition per se that accounts for deprenyl's neuroprotective effect (Finnegan 1993; Finnegan et al. 1990). Perhaps it is the propargylamine moiety of deprenyl that is responsible for its inhibition of DSP-4 neurotoxicity (Maruyama et al. 2000).

## 5.4.3 Specificity of DSP-4

Generally, because DSP-4 is preferred over xylamine, the latter is used in few studies today. Consequently, most of the following discussion relates to DSP-4.

Because of its high affinity for the NET, DSP-4 is selective for noradrenergic neurons and has little direct effect on dopaminergic, cholinergic, GABAergic, glutamatergic, and glycinergic neurons (see Dudley et al. 1990; Jaim-Etcheverry and Zieher 1983). In the brain, DSP-4 has greatest neurotoxic effects on noradrenergic nerve terminals of locus coeruleus origin (Fritschy and Grzanna 1989, 1991a). DSP-4 degenerative effects in the brain are long-lived. In contrast, in the periphery, sympathetic noradrenergic neurons recover from DSP-4 effects often in less than 1 week (Jaim-Etcheverry and Zieher 1980; Jonsson et al. 1981). Serotoninergic nerves are the only other monoaminergic nerve prominently affected by DSP-4, and the effect on serotonin (5-HT) content and serotoninergic innervation of the brain is long-lived (Jonsson et al. 1981).

## 5.4.4 Long-Lived Effects of DSP-4 Treatment

When adult rats are treated with a high dose of DSP-4, noradrenergic innervation of projection areas of the locus coeruleus (i.e., neocortex, hippocampus, cerebellum, spinal cord) is markedly reduced, and the effect is long-lived (Jaim-Etcheverry and Zieher 1980; Jonsson et al. 1982). Regions of the brain that are innervated by noradrenergic fibers originating from non-coerulear nuclei are little affected by DSP-4.

When neonatal rats are treated with a high dose of DSP-4, noradrenergic innervation of projection areas of the locus coeruleus is markedly reduced, as is observed in rats treated as adults. In contrast, regions of the brain proximal to the locus coeruleus (i.e., cerebellum, pons, medulla) become hyperinnervated by noradrenergic neurons, and regional NE content is thus elevated. Hyperinnervation is the consequence of collateral axonal fiber sprouting, a reactive sprouting attributable in large part to the degeneration of distal projection of locus coeruleus axons to the neocortex, hippocampus, and spinal cord. This effect is analogous to that seen after surgical transection of the dorsal bundle, the axonal tract projecting from locus coeruleus to forebrain (Kostrzewa et al. 1988). Regions of the brain innervated by non-coerulear nuclei are little affected by DSP-4, as are the sympathetic noradrenergic fibers in the periphery (Bortel et al. 2008; Jaim-Etcheverry and Zieher 1980; Jonsson et al. 1982; Nowak et al. 2006).

## 5.4.5 Summary on DSP-4

Although 6-OHDA is the most frequently used neurotoxin for noradrenergic neurons, there are occasional studies in which DSP-4 is advantageous. This relates to DSP-4 (1) relative selectivity, particularly for locus coeruleus neurons;

also (2) there is the outcome of long-lived noradrenergic terminal denervation and (3) lack of effect on dopaminergic neuronal fields. Additionally, (4) DSP-4 is able to cross the blood-brain barrier, and thus DSP-4 can be administered in the periphery to produced CNS effects in adults, unlike 6-OHDA. (5) When stored, DSP-4 is more stable than 6-OHDA.

DSP-4 and xylamine neurotoxicity are discussed in greater detail in several reviews (Bortel 2013 [in this *Handbook of Neurotoxicity*]; Dudley et al. 1990; Jaim-Etcheverry 1998; Jaim-Etcheverry and Zieher 1983).

## 5.5 1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP)

## 5.5.1 Historical Perspective

In 1976, a graduate student chemist in Maryland synthesized the meperidine analog 1-methyl-4-phenyl-4-propionoxypiperidine (MPPP), to be used as an opioid substance of abuse. However, during the process of synthesis, the chemical mixture was heated to too high a temperature, resulting in the formation of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP). This contaminant, 1-3 % of the MPPP product, proved to be neurotoxic to dopaminergic neuronal perikarya in the pars compacta substantia nigra (SNpc). Within days, he developed the permanent symptomology spectrum of Parkinson's disease (PD) (see Shafer 1985).

In 1979 in Southern California, a similar synthetic street drug, termed "China White" – the moniker for the purest Southeast Asia heroin – produced the same PD symptomology in substance abusers, again a permanent effect. Three years later, the same event occurred in Northern California, resulting in multiple young substance abusers developing PD-like symptoms (lifelong) and death to more than 50 people. The neurologists J. William Langston and Philip Ballard in San Jose recognized that a bad batch of a heroin-like drug was responsible, and they succeeded in alerting the public to the link between China White and PD/ death. As a consequence, that street drug disappeared and the appearance of new cases of early-age PD abated (Langston and Ballard, 1984; Ballard et al. 1985).

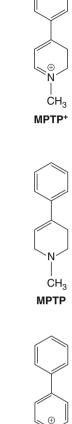
Subsequently, MPTP was identified as the neurotoxic contaminant in China White, as testing in animals replicated the forensic finding of SNpc cell destruction found in China White substance abusers. Ultimately, MPTP became the prime neurotoxin to replicate PD in animals. MPTP is of value for probing cellular mechanisms and processes associated with PD and in studying neuroprotectants and new treatment approaches (Shafer 1985; Pasquali et al. 2013).

The key terms "MPTP" or "1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine" appear in 5,746 papers in PubMed, as of January 30, 2013. Key terms for the MPTP metabolite neurotoxic species "1-methyl-4-phenyl-1,2,3,6-tetrahydropyridinium" or "MPP<sup>+</sup>" appear in 2,664 papers in PubMed as of January 30, 2013.

## 5.5.2 MPTP Mechanism and Selectivity of Action

MPTP (Fig. 8) crosses the blood-brain barrier (BBB) (Riachi et al. 1988) and is converted by monoamine oxidase-B (MAO-B) in astrocytes (Chiba et al. 1984;

Fig. 8 (From Kostrzewa 1999b)



CH<sub>3</sub>

MPP<sup>+</sup>

Fig. 9 (From Kostrzewa 1999b)

**Fig. 10** (From Kostrzewa 1999b)

Westlund et al. 1985; Di Monte et al. 1991) [and to a lesser extent by MAO-A in other cells (Singer et al. 1986)] to the intermediate 1-methyl-4-phenyl-1,2,3-dihydropyridinium ion (MPDP<sup>+</sup>) (Fig. 9) which undergoes spontaneous twoelectron oxidation to 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridinium ion (MPP<sup>+</sup>) (Fig. 10) (Castagnoli et al. 1985). MPP<sup>+</sup>, released from astrocytes and having high affinity for the DAT, is actively accumulated by dopaminergic neurons (Chiba et al. 1985; Javitch and Snyder 1984; Javitch et al. 1985). Inside DA nerves, the mitochondrial membrane potential actively drives accumulation of MPP<sup>+</sup> to a level in mitochondria as much as 40 times the concentration in cytoplasm (Wu et al. 1990). Here, MPP<sup>+</sup> inhibits complex I in the respiratory transport chain, between NADH dehydrogenase and coenzyme Q, thereby preventing ATP formation (Chan et al. 1993) and promoting formation of reactive oxygen species (ROS). MPP<sup>+</sup> additionally inhibits alpha-ketoglutarate dehydrogenase in the tricarboxylic acid cycle, thereby preventing succinate formation, a substrate for complex II (Mizuno et al. 1987). Cellular energy depletion (Di Monte et al. 1986; Denton and Howard 1987) and ROS damage to critical elements result in neuronal damage and subsequent neuronal degeneration (Ramsay et al. 1991; see Pasquali et al. 2013). Glial-derived inflammatory cytokines add to stresses on the neuron (see Teismann et al. 2003). DA innervation of neostriatum and DA neuronal perikarya in SNpc are largely affected, although DA neurons in the ventral tegmental nucleus (VTN) and noradrenergic (melanized) neurons in locus coeruleus also are damaged (Seniuk et al. 1990).

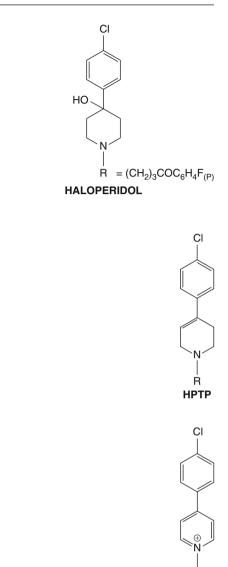
## 5.5.3 Interspecies Effects of MPTP

C57 BL/6 black mice are more sensitive than other mouse strains to MPTP (Sonsalla and Heikkila 1988; Muthane et al. 1994), while rats are largely resistant to MPTP partly because of the abundance of MAO-B in the cerebrocapillary endothelium which constitutes the BBB. MPTP in rats is thereby converted to MPP<sup>+</sup> before crossing the BBB; MPP<sup>+</sup>, being ionized, is unable to cross the BBB to enter the brain (Kalaria et al. 1987). Additionally, DA neurons in rats are imbued with a greater abundance of vesicular monoamine transporter-2 (VMAT-2) versus mice. This facet relates to a greater reservoir of MPP<sup>+</sup> inside the vesicle, which translates to a lower cytoplasmic concentration of MPP<sup>+</sup> in rats versus mice (Staal and Sonsalla 2000). This too accounts for a lowered neurotoxic potential for MPP<sup>+</sup> in rats. Nonhuman primates are highly susceptible to MPTP neurotoxicity, with MPTP resulting in effects similar to that observed in human parkinsonian patients (Jenner et al. 1984). It has become conventional to produce unilateral MPTP lesioning in nonhuman primates (e.g., intracarotid administration on one side), so that animals are able to remain mobile and eat without assistance (Bankiewicz et al. 1986; see Fox and Brotchie 2010).

Royland and Langston (1998), Teismann et al. (2003), as well as Carta et al. (2013) and Pasquali et al. (2013) in this *Handbook of Neurotoxicity* provide detailed discussion and analysis of MPTP neurotoxicity mechanisms and MPTP modeling of Parkinson's disease in rodents and nonhuman primates.

## 5.6 Haloperidol-Related Chemical Species

The classical antipsychotic drug haloperidol (Fig. 11), a pyridinium chemically similar in structure to MPTP, is regarded as a neurotoxin or protoxin (Roberts et al. 1995). Haloperidol and its dehydration product 4-(4-chlorophenyl)-1,4-(4-fluorophenyl)-4-oxobutyl-1,2,3,6-tetrahydropyridine (HPTP) (Fig. 12) are oxidatively metabolized to HPP<sup>+</sup> (Fig. 13) and 4-(4-chlorophenyl)-1,4-(4-fluorophenyl)-4-hydroxybutylpyridinium (RHPP<sup>+</sup>) (Subramanyam et al. 1991; Eyles et al. 1994, 1996; Igarashi et al. 1995; Usuki et al. 1996) – analogs of MPTP. In vitro HPP<sup>+</sup> is **Fig. 11** (From Kostrzewa 1999b)





1999b)

Fig. 12 (From Kostrzewa

**Fig. 13** (From Kostrzewa 1999b)

neurotoxic to both mesencephalic cells (Bloomquist et al. 1994) and neuroblastoma cells (Fang and Yu 1995). In vivo HPP<sup>+</sup> is an inhibitor of the mitochondrial respiratory chain (Rollema et al. 1994) and overtly neurotoxic to DA nerves (Subramanyam et al. 1990). It is these chemical derivatives that perhaps account for the overt neuronal destruction (Benes et al. 1984, 1985a, b) of gamma-aminobutyric acid (GABA) neurons, glutamate neurons (Meshul and Casey 1989;

R

HPP<sup>+</sup>

**Fig. 14** (From Kostrzewa 1999b)

Meshul et al. 1994), and cholinergic neurons (Mahadik et al. 1988) associated with long-term haloperidol treatment of schizophrenics, in whom tardive dyskinesia has developed. The neurological aspect of haloperidol in tardive dyskinesia is developed further in the *Handbook of Neurotoxicity* chapter on tardive dyskinesia (Kostrzewa et al. 2013).

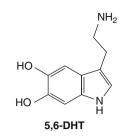
# 5.7 2'-Amino-MPTP

2'-Amino-MPTP (2'-NH<sub>2</sub>-MPTP), unlike the parent MPTP, had virtually no effect on dopaminergic neurons but did effectively reduce striatal 5-HT and 5-HIAA (Johannessen et al. 1987). Ultimately 2'-NH<sub>2</sub>-MPTP was found to also produce a long-term reduction of NE as well as 5-HT contents in rodent forebrain (Andrews and Murphy 1993a, b; Luellen et al. 2003). A substrate for both MAO-A and MAO-B, 2'-NH<sub>2</sub>-MPTP likely exerts its effects via the metabolite 2'-NH<sub>2</sub>-MPP<sup>+</sup> which has affinity to the NET and SERT. Effects of 2'-NH<sub>2</sub>-MPTP on NE and 5-HT in the brain are blocked by desigramine and fluoxetine, respectively (Andrews and Murphy 1993c). Intracellular 2'-NH<sub>2</sub>-MPP<sup>+</sup>, like MPP<sup>+</sup>, apparently acts on mitochondria and other cellular elements to form superoxide. Mice overexpressing Cu-Zn superoxide dismutase, which converts superoxide to less reactive H<sub>2</sub>O<sub>2</sub>, are protected from the effects of 2'-NH<sub>2</sub>-MPP<sup>+</sup> (Andrews et al. 1996). Unlike MPP<sup>+</sup>, 2'-NH<sub>2</sub>-MPP<sup>+</sup> is a poor substrate for VMAT-2 (Numis et al. 2004).

The neurotoxicity of 2'-NH<sub>2</sub>-MPTP is discussed in detail in this *Handbook of Neurotoxicity* by Ochroch et al. (2013).

## 5.8 5,6- and 5,7-Dihydroxytryptamine (5,6-DHT, 5,7-DHT)

5,6-Dihydroxytryptamine (5,6-DHT) (Fig. 14), a di-/ortho-phenolic analog of 5-hydroxytryptamine (5-HT, serotonin) (Fig. 15), and 5,7-dihydroxytryptamine (5,7-DHT) (Fig. 16), a di-/meta-phenolic analog of 5-HT, were characterized as serotoninergic neurotoxins by Hans Baumgarten in early 1970s – shortly after the discovery of 6-hydroxydopamine (6-OHDA) as a neurotoxin for noradrenergic and dopaminergic neurons (Baumgarten et al. 1971, 1972a, b; Baumgarten and Schlossberger 1973; Baumgarten and Lachenmeyer 1972).



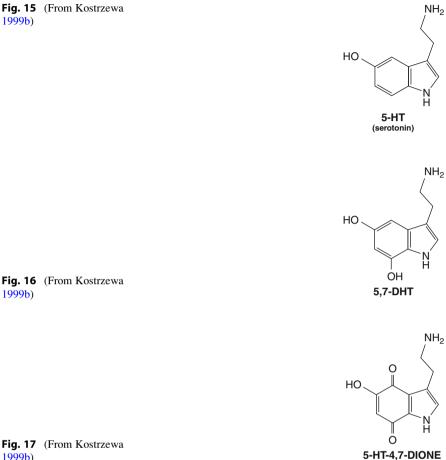
1999b)

Fig. 16 (From Kostrzewa 1999b)

1999b)

#### 5.8.1 Mechanisms of Action of 5,6- and 5,7-DHT

5.6- and 5.7-DHT owe their high degree of selective serotoninergic neurotoxicity to their high affinity to the serotonin transporter (SERT) (Björklund et al. 1973a, b, 1975b), which selectively transports the DHTs into the 5-HT neuron. Once present in high intraneuronal concentration, 5,6- and 5,7-DHT intracyclize to generate quinones and additional putative neurotoxic indolamine semiquinoidal and quinoidal species, with 5-HT-4,7-dione (Fig. 17) and its dimer 6,6'-bi(5-HT-4,7dione) being the principle derivatives (Tabatabaie and Dryhurst 1992, 1998). All are highly reactive. The resulting formation of cellular ROS (e.g., hydrogen peroxide,  $H_2O_2$ ; superoxide,  $O_2^-$ ; hydroxyl radical, HO<sup>•</sup>) further burdens the protective capacity of 5-HT neurons. Ultimate damage of mitochondrial cytochromes (Cohen and Heikkila 1978), cellular lipids, and cellular proteins thus compromises function, resulting in cellular destruction (Baumgarten et al. 1978a, 1981, 1982; Tabatabaie and Dryhurst 1992; Tabatabaie et al. 1990, 1993).



5,6- and 5,7-DHT neurotoxicity are discussed in greater detail in several reviews (Baumgarten et al. 1973, 1978a, b; Tabatabaie and Dryhurst 1998; Pranzatelli 1998; Baumgarten and Lachenmayer 2004) and in a later chapter of this *Handbook of Neurotoxicity* (Paterak and Stefański 2013).

## 5.8.2 Limitations of 5,6-DHT

Although the biological effects of 5,6-DHT are similar to those of 5,7-DHT, higher doses of 5,6-DHT become less selective for 5-HT neurons and produce convulsions and death (Baumgarten and Lachenmeyer 1972; Baumgarten et al. 1972b, 1973; Björklund et al. 1973b). Consequently, 5,7-DHT has supplanted 5,6-DHT as a serotoninergic neurotoxin.

## 5.8.3 Effects of 5,7-DHT in Adults

**Long-Lived Effects in the Brain of Animals Lesioned as Adults with 5,7-DHT** Although 5,7-DHT is relatively selective for serotoninergic neurons, a low dose of 5,7-DHT will nonetheless produce both 5-HT and NE depletion in the brain (Baumgarten et al. 1973). The effects of 5,7-DHT on NE neurons can be largely prevented by pretreatment with the NET inhibitor desipramine (Björklund et al. 1975a).

A high dose of 5,7-DHT will destroy 5-HT innervation in virtually all regions of the CNS. The pons and medulla – the brain regions nearest to the most prominent 5-HT perikarya, the dorsal and midbrain raphe nuclei – are least affected by 5,7-DHT treatment, as 5-HT content is only modestly altered after 5,7-DHT. In contrast, regions of the CNS most distal to the dorsal and medial raphe are prominently affected by 5,7-DHT treatment, as 5-HT content of the neocortex, hippocampus, and spinal cord is depleted about 85–90 %. Serotoninergic denervation of striatum by 5,7-DHT is profound. These 5,7-DHT-induced alterations in 5-HT innervation in the brain are long-lived (Baumgarten et al. 1973).

# Long-Lived Effects in the Brain of Animals Lesioned as Neonates with 5,7-DHT

When administered to neonatal rats, a single high dose of 5,7-DHT produces lifelong, marked (~90 %) 5-HT denervation and 5-HT depletion of the neocortex, hippocampus, and striatum (Jonsson et al. 1978). In neonatally 5,7-DHT-lesioned rats studied in adulthood, there is prominent DA D<sub>2</sub>-receptor supersensitivity, as evidenced by enhanced behavioral responses to the D<sub>2</sub> agonist quinpirole (Brus et al. 1995). In these neonatally 5,7-DHT-lesioned rats, there is also a reduction in histamine levels in the brain and enhanced histamine H<sub>3</sub> receptor antagonist-evoked oral and locomotor activity – indicative of alterations in histaminergic neuronal systems and their effects (Jośko et al. 2011).

In rats co-lesioned as neonates with 5,7-DHT and the noradrenergic neurotoxin DSP4, prominent  $D_2$ -receptor supersensitivity is evident in adulthood (Nowak et al. 2009).

In rats lesioned neonatally with 6-OHDA, there is lifelong supersensitization of DA  $D_1$  receptors, as evidenced by enhanced locomotor and oral activity responses and stereotypic responses to the  $D_1$  agonist SKF 38393 (Kostrzewa and Gong 1991;

Gong et al. 1993). However, when 5,7-DHT is administered neonatally, simultaneous with 6-OHDA treatment, there is long-term (into adulthood) suppression of  $D_1$ -receptor supersensitivity (Brus et al. 1994). Spontaneous locomotor activity is normally increased in adulthood rats that were lesioned as neonates with 6-OHDA; and there is even greater spontaneous locomotor activity in those rats simultaneously lesioned as neonates with 6-OHDA and 5,7-DHT. Rats so-lesioned as neonates with 6-OHDA and 5,7-DHT have been proposed as an animal model of attention deficit hyperactivity disorder (ADHD) (Kostrzewa et al. 1994).

## Summary on 5,7-DHT

5,7-DHT, a relatively selective neurotoxin for 5-HT neurons, provides a means to reasonably study acute and long-lived effects of 5-HT denervation – in both neonatal and adult animals. 5,7-DHT likewise is useful for determining the influence of prominent 5-HT denervation on the effects of other neuronal phenotypic systems and in elucidating the role of 5-HT per se in a variety of functions, including behaviors.

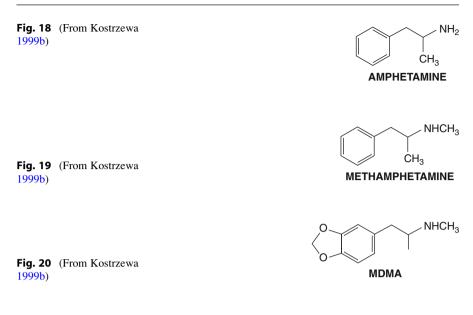
# 5.9 Amphetamines

## 5.9.1 Amphetamine (AMPH)

Amphetamine (AMPH) (Fig. 18), chemically similar to DA and NE, has high affinity for the DAT, NET and also for the serotonin transporter, SERT. By virtue of these effects, AMPH gains entry into DA-, NE-, and 5-HT neurons. Intraneuronal AMPH accumulates in both cytoplasm and vesicles, displacing these monoamines and reversing the transporters to evoke monoamine release. In additon, as a consequence of AMPH competition with DA and NE for the DAT and NET, respectively, extraneuronal and synaptic effects of both DA and NE are prolonged and enhanced. Nevertheless, acute treatment or intermittent AMPH treatments are not neurotoxic (Westfall and Westfall 2006).

However, when continuously administered for a prolonged period ( $\geq$ 3d), AMPH becomes overtly neurotoxic to striatal dopaminergic nerves, as evidenced by long-term striatal DA depletion (Wagner et al. 1980), loss of tyrosine hydroxylase (TOH) activity (Ellison et al. 1978), loss of DAT number and DAT protein (Scheffel et al. 1996; Krasnova et al. 2001), presence of swollen histofluorescence axons, reduced numbers of dopaminergic terminals, and overt indices of neurodegeneration – deposition of argyrophilic granules (silver degeneration staining) and ultrastructural evidence of terminal damage (Ellison et al. 1978; Nwanze and Jonsson 1981; Ryan et al. 1988, 1990; Ellison 1992). AMPH neurotoxicity extends to TOH-positive mesencephalic neurons and cortical cells (Stumm et al. 1999; Jakab and Bowyer 2002).

AMPH neurotoxicity is associated with hyperthermia (Freh 1975; Clausing and Bowyer 1999) which magnifies the effect of AMPH-induced production of superoxide and hydroxyl radical (Lin et al. 1991; Kil et al. 1996; Huang et al. 1997a; Krasnova et al. 2001) as well as DA-quinone (LaVoie and Hastings 1999).

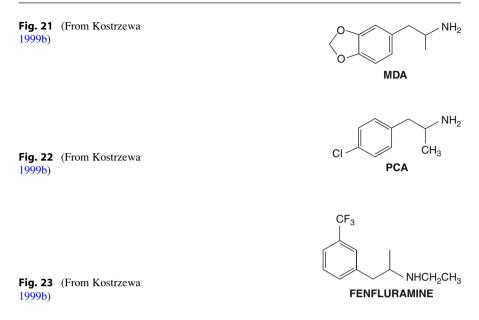


# 5.9.2 Methamphetamine (METH, Speed, Crank)

The overall acute effects of methamphetamine (METH) (Fig. 19) are analogous to those of AMPH (see Krasnova and Cadet 2009). Likewise, neurotoxic effects are similarly related to generation of ROS including DA-quinones (LaVoie and Hastings 1999), hydrogen peroxide, and superoxide (Stokes et al. 1999; Miyazaki et al. 2006). However, when METH was added to an in vivo microdialysis perfusate into striatum, the hydroxyl radical level was not elevated (Pereira et al. 2004). It appears that METH-induced glutamate exocytosis (Marshall et al. 1993; Abekawa et al. 1994; Mark et al. 2004) may be largely responsible for generation of nitric oxide (NO) and other ROS (Itzhak et al. 1998; Imam et al. 2001; Itzhak and Ali 2006). Serotoninergic neuronal degeneration is prominent with METH (Ricaurte et al. 1982). The DA D<sub>1</sub>-receptor antagonist SCH 23390 [R-(+)-7-chloro-8-hydroxy-3-methyl-1-phenyl-2,3,4,5-tetrahydro-1H-3-benzazepine] attenuates neurotoxic effects of METH on DA and 5-HT systems, while the DA D<sub>2</sub> antagonist sulpiride attenuates METH neurotoxic effects on DA systems (Sonsalla et al. 1986).

# 5.9.3 Methylenedioxymethamphetamine (MDMA, Ecstasy)

Methylenedioxymethamphetamine (MDMA) (Fig. 20), a substrate for the 5-HT transporter (SERT), is accumulated by 5-HT nerves; then MDMA is accumulated by vesicles via the vesicular monoamine transporter (VMAT), whereby it releases 5-HT from vesicles and into the synapse (Baumann and Rothman 2009). This action ultimately results in DA exocytosis. MDMA is selectively neurotoxic to 5-HT terminals in rats and nonhuman primates (Green et al. 1995). In mice, MDMA damages nigrostriatal dopaminergic fibers (Commins et al. 1987; Cadet et al. 1995). Methylenedioxyamphetamine (MDA) (Fig. 21) acts similarly to MDMA.



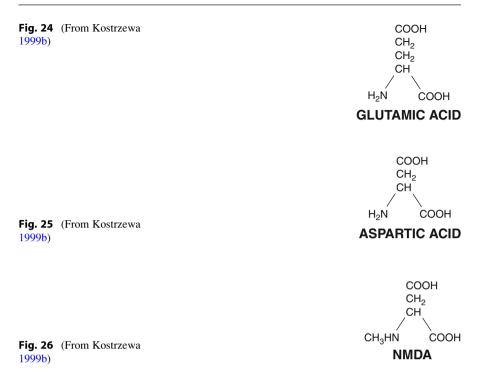
## 5.9.4 Parachloroamphetamine (PCA)

Parachloroamphetamine (PCA) (Fig. 22) was initially found to produce longterm reduction in tryptophan hydroxylase (Trp-OH) activity and 5-HT in the rat brain (Sanders-Bush et al. 1972a, b). Silver staining later established that PCA actions were related to overt neurotoxic effects, primarily on the serotoninergic B9 cell group (Harvey et al. 1975). PCA neurotoxicity is dependent on 5-HT release; prior 5-HT depletion with parachlorophenylalanine and reserpine prevents PCA neurotoxic actions (Berger et al. 1992). However, prior destruction of striatal dopaminergic innervation does not negate the neurotoxic effect of PCA (Perry et al. 1995).

AMPH, METH, MDA, and analogs have been the subject of many reviews (Cadet et al. 2007; Yamamoto and Raudensky 2008; Gouzoulis-Mayfrank and Daumann 2009; Yamamoto et al. 2010; Steinkellner et al. 2011) including the chapter by Bisagno and Cadet (2013) in this *Handbook of Neurotoxicity*.

## 5.9.5 Fenfluramine

Another analog of amphetamine, namely, fenfluramine (Fig. 23), also is neurotoxic towards serotoninergic fibers. Fenfluramine, in high dose or in repeated doses, depletes 5-HT in brain (Harvey and McMaster 1975; Steranka and Sanders-Bush 1979; Schuster et al. 1986), also reduces tryptophan hydroxylase, the rate-limiting enzyme in 5-HT synthesis (Steranka and Sanders-Bush 1979), additionally reduces SERT number (Schuster et al. 1986; Appel et al. 1990; Zaczek et al. 1990) and reduces the numbers of 5-HT immunoreactive fibers in the brain (Appel et al. 1989; Molliver and Molliver 1990). The combination of phentermine with fenfluramine as an anorectic agent was withdrawn from the market because of the association with valvular heart disease (Connolly et al. 1997).



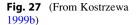
# 5.10 Excitatory Amino Acids

# 5.10.1 Historical Perspective

In 1957, Lucas and Newhouse (1957) reported that iontophoretic application of glutamic acid (GLU) (Fig. 24) produced overt destruction of the inner layers of immature mouse retina. Curtis et al. (1959) next showed that GLU and aspartic acid (ASP) (Fig. 25) are excitatory (i.e., produce depolarization) to spinal neurons. And Olney and colleagues (Olney 1969; Olney and Sharpe 1969) demonstrated that GLU is specifically neurotoxic in mice and monkeys, giving birth to the concept of "excitotoxicity" and the range of action of excitatory amino acids (EAAs) (Olney et al. 1971).

# 5.10.2 Mechanisms of GLU Excitotoxicity

GLU-mediated neurotoxicity is related to its agonist action at NMDA (N-methylaspartate) (Fig. 26) receptors (NMDA-R), kainate-R (Fig. 27), or AMPA-R (AMPA = alpha-amino-2,3-hydroxy-5-methyl-3-oxo-isoxazole-propionic acid) (Fig. 28), resulting in massive influx of Na<sup>+</sup>, Cl<sup>-</sup>, and Ca<sup>2+</sup> into postsynaptic neurons (Rothman 1985; Choi 1987) and delayed activation of Ca<sup>2+</sup>-sensitive proteases (Frandsen and Schousboe 1992; Tymianski et al. 1993). As a consequence of these actions and in concert with release of intracellular Ca<sup>2+</sup> stores, ultimate dendrosomal swelling with vacuolization and mitochondrial edema of postsynaptic neuronal elements concludes in neuronal cell death (see Olney 1981;



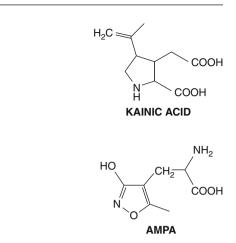


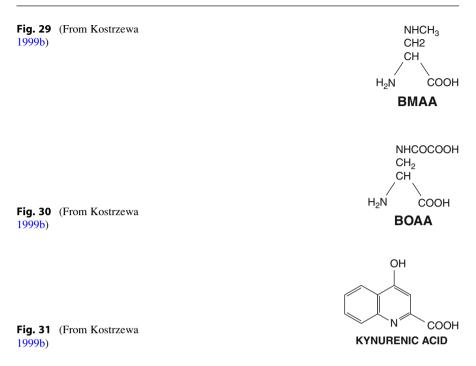
Fig. 28 (From Kostrzewa 1999b)

Mody and MacDonald 1995). A variety of intracellular proteins, when activated by ionic events, play a major role in cellular events relating to metabolic inhibition, apoptosis, or necrotic cellular demise (Zeevalk and Nicklas 1991; Ikonomidou and Turski 1996).

As a consequence of the above cellular events, cellular ROS are generated, and these add to the challenge of cellular viability. Neuronal  $O_2^{-}$  is converted by superoxide dismutase to oxygen ( $O_2$ ) and hydrogen peroxide ( $H_2O_2$ ). Neuronal nitric oxide synthase (nNOS) activation leads to cellular NO formation (Dawson et al. 1996). In turn, NO reacts with  $O_2^{-}$ , forming peroxynitrite (ONOO<sup>-</sup>) (Huie and Padmaja 1993), a highly reactive species that produces protein nitration, lipid peroxidation, impairment of GAPDH, inhibition of complexes I and II of the electron transport chain, and DNA injury (Radi et al. 1991, 1994; van der Vliet et al. 1998; Hara et al. 2005).

# **5.10.3 Excitotoxicity in Neurological and Neurodegenerative Disorders** Ischemia, Anoxia, and Traumatic Brain Injury

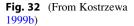
During transient hypoxia, there is impaired uptake of GLU and ASP from extracellular spaces, leading to a 5-fold or greater increase in GLU and ASP extracellular levels (Beneviste et al. 1984; Globus et al. 1988). This is accompanied by neurotoxicity in cerebral cortex (Jorgensen and Diemer 1982), in hippocampal CA1 pyramidal neurons, and in striatal medium size neurons (Rothman and Olney 1986). Reperfusion results in augmented release of GLU and ASP (Silverstein et al. 1986), resulting in greater excitotoxicity. Prior lesioning of the afferent glutamatergic pathway prevents neurodegeneration during ischemia (Wieloch et al. 1985), as does NMDA-R antagonist treatment (Pulsinelli et al. 1993; Waner et al. 1995) including group I metabotropic GLU-R antagonists (Kingston et al. 1999). In traumatic brain injury (TBI), similarly there is an increase in extracellular glutamate (Faden et al. 1989; Nilsson et al. 1990) accompanied by neurodegeneration, with neuroprotection conferred by NMDA-R antagonists (McIntosh et al. 1989).



### **Neurodegenerative Disorders**

The involvement of glutamatergic systems in the neurogenesis of PD is indicated by the findings that MPP<sup>+</sup>-induced neurodegeneration is attenuated by NMDA-R antagonists (Turski et al. 1991; Santiago et al. 1992). The naturally occurring excitatory amino acids  $\beta$ -N-methylamino-L-alanine (L-BMAA) (Fig. 29) and  $\beta$ -Noxalylamino-L-alanine (L-BOAA) (Fig. 30) are linked, respectively, to Guam ALS-parkinsonism dementia syndrome (Spencer 1987) and neurolathyrism (Spencer et al. 1986). The kainic acid analog domoic acid, identified as the active principal in contaminated mussels, was responsible for an outbreak of toxic encephalopathy in Canada, resulting in motor neuronopathy and partial paralysis (Perl et al. 1990; see Pérez-Gómez and Tasker 2013). Kainate per se is associated with degeneration of GABA- and substance P-containing neurons (Coyle and Schwarcz 1976; McGeer and McGeer 1976).

Quinolinic acid produces a striatal lesion the models Huntington's disease (HD), associated with chorea in humans (Schwarcz et al. 1983; Beal et al. 1986), although quinolate levels are not elevated in HD (Reynolds et al. 1988; Heyes et al. 1991). Rather, animal modeling of HD is best replicated by aminooxyacetic acid (AOAA) (Urbanska et al. 1989, 1991), a synthesis inhibitor of the endogenous EAA antagonist kynurenic acid (Fig. 31) (Turski et al. 1989; see Urbanska et al. 2013). Moreover, AOAA neurotoxicity is attenuated by NMDA-R antagonists (Urbanska et al. 1991). In HD, kynurenine aminotransferases – enzymes forming kynurenine – are deficient in basal ganglia, thereby implicating dysfunctional kynurenine synthesis in the genesis of HD (Jauch et al. 1995). The overt mechanism of AOAA





neurotoxicity relates to impaired cellular metabolism (Beal et al. 1991), and the mitochondrial toxin 3-nitropropionic acid (3-NP) is additionally engaged in rodents and primates to replicate the lesions attending HD (Palfi et al. 1996), with this neurotoxicity likewise being attenuated by NMDA-R antagonists (Brouillet et al. 1995).

The accumulation of extraneuronal beta-amyloid in Alzheimer disease (AD) is an incipient contributor to the neuropathology of this disorder. Beta-amyloid promotes microglial exocytosis of an unknown NMDA-R agonist (Giulian et al. 1995), stimulates NO production (Goodwin et al. 1995), enhances GLU excitotoxicity (Arias et al. 1995), impairs GLU uptake by astrocytes (Harris et al. 1996), and destabilizes intraneuronal  $Ca^{2+}$  (Koh et al. 1990).

In amyotrophic lateral sclerosis (ALS), there is an apparent loss of glial EAAT-2 glutamate transporter in the brain and spinal cord (Rothstein et al. 1992, 1995), and this perhaps accounts for elevated CSF levels of GLU in a population of ALS patients (Shaw et al. 1995). The CSF of ALS patients was found to be neurotoxic to neurons in culture, by virtue of kainate-R and AMPA-R action (Couratier et al. 1993, 1994). Motor neurons in ALS display atypical Ca<sup>2+</sup>-permeable AMPA-R, conferring selective vulnerability to GLU excitotoxic action (Williams et al. 1997).

Comprehensive and insightful reviews have been written on the topic of glutamate, excitatory amino acids, and excitotoxicity (Urbanska et al. 1998; Turski and Ikonomidou 2013).

## 5.11 NMDA Receptor Antagonists

While NMDA-R antagonists are known to attenuate the neurotoxic effects of NMDA-R agonists (Wieloch 1985; Weiss et al. 1986; Choi et al. 1988; Park et al. 1988; Clifford et al. 1990), paradoxically NMDA-R antagonists are associated with neurotoxicity – according to the dosing schedule and stage of ontogeny at the time of treatment (Olney et al. 1989; Fix et al. 1995, 1996; Farber et al. 1995; Wozniak et al. 1996). Also, noncompetitive NMDA-R antagonists exact a greater toll than competitive NMDA-R antagonists (Ellison 1994). Dizocilpine (MK-801), ketamine, and phencyclidine (PCP) (Fig. 32) produce neurotoxicity to pyramidal neurons in cortical layers II and IV of posterior cingulate/retrosplenial cortex (Olney et al. 1989), also in striatum, subiculum and hippocampal areas CA1 and CA3 (Fix et al. 1995), cingulate cortex, fasciculus retroflexus (FR), and lateral habenula (Ellison 1995).

As proposed by Olney et al. (1991), NMDA-R antagonists disinhibit GABA neurons, which results in excess cholinergic activity. Glycine agonists and polyamine agonists attenuate the toxicity of NMDA-R antagonists without themselves causing neurotoxicity (Duval et al. 1992; Hargreaves et al. 1993). This hypothesis is based on the findings that NMDA-R neurotoxicity is prevented by muscarinic-,  $\sigma$ -, and  $\alpha_2$ -R antagonists and both typical and atypical antipsychotics (Olney et al. 1991; Farber et al. 1993, 1995).

NMDA-R neurotoxicity has been the topic of several reviews (Olney et al. 1991; Rajdev and Sharp 1998; Olney 2002; Low and Roland 2004).

# 5.12 Cocaine

Neurotoxic effects of the abused drug cocaine have come to light, largely in the past two decades. Cocaine is known to impair mitochondrial complex I, II, and III activity and reduce state 3 respiration (Devi and Chan 1997; Yuan and Acosta 2000; Dietrich et al. 2004). This is followed by loss of the mitochondrial membrane potential and release of cytochrome c (see Cunha-Oliveira et al. 2008). At the neurotoxic stage, changes occur in Bcl-2 and Bax (Dey et al. 2007). Apoptosis is the outcome (see Cunha-Oliveira et al. 2008).

Prolonged (5-day) continuous administration of cocaine to adult rats via a cocaine-silicone pellet produced degeneration of the lateral habenula (LHb) in the dorsal thalamus and its projection – the fasciculus retroflexus (FR) – to midbrain dopaminergic nuclei (substantia nigra, ventral tegmental nucleus) and serotoninergic nuclei (dorsal and medial raphe) (Ellison 1992). AMPH, METH, MDMA, and cathinone were later found to produce similar but not identical effects as cocaine (Ellison 2002). Also, when cocaine was administered to pregnant rats for the remainder of pregnancy, starting on gestation day 17, the LHb and FR of rat pups underwent analogous neurodegeneration (Murphy et al. 1999).

The neurotoxicity of prolonged administration of cocaine and related agents (e.g., AMPH) resembles that of phencyclidine (Ellison 1995) and is thought to be related to a reduction in glucose metabolism in LHb (London et al. 1986; Wirtshafter et al. 1994), consequent to loss of GABA inhibition and associated excitotoxic neurotoxicity (Ellison 1998).

The neurotoxic effects of relatively short-term cocaine administration are discussed in papers by Cunha-Oliveira et al. (2008, 2013). The neurotoxic effects of continuous and prolonged cocaine administration are discussed by Ellison (1998).

## 5.13 AF64A (Ethylcholine Aziridinium Ion)

AF64A is a Trojan horse neurotoxin with relative selectivity for cholinergic nerves. Chemically similar to choline but with an aziridinium moiety on the nitrogen atom, AF64A is a nitrogen mustard (alkylating agent) that targets the choline uptake site on acetylcholine (Ach)-containing nerves (Fisher and Hanin 1986, Hanin 1988, 1996). In high concentration (>22  $\mu$ M), AF64A alkylates both low- and high-affinity choline transporter sites, impairing its own uptake by the nerve and producing nonspecific damage to any tissue but notably to the cerebral endothelium which maintains a choline transport system (Gomez-Ramos et al. 1990).

In low concentration ( $<5 \mu$ M), AF64A is reversibly bound to the choline transporter and then largely enters the cholinergic nerve and alkylates many sites, predominately enzymes and proteins with affinity for choline: choline acetyltransferase, choline kinase, acetylcholinesterase, and choline dehydrogenase (Curti and Marchbanks 1984; Hortnagl and Hanin 1992; Hortnagl et al. 1988; Rylett and Colhoun 1984; Uney and Marchbanks 1987). Alkylated enzymes and other macromolecules become inactivated, rendering the nerve dysfunctional (Barlow and Marchbanks 1984; Leventer et al. 1987; Potter et al. 1987; Sandberg et al. 1985). Cell death is related to alkylation at N-7 guanine sites on DNA, resulting in strand breaks and termination of RNA transcription (Futscher et al. 1992). AF641-induced alterations in cholinergic nerves are often delayed by as much as 48 h (Hanin 1996).

Unable to cross the blood-brain barrier, AF64A is commonly administered into the cerebral ventricles to model for Alzheimer's type senile dementia (Fisher and Hanin 1986). Although the levels of neurotransmitters other than Ach are acutely altered by AF64A, these changes are short-lived and thus considered to be secondary to the abrupt alteration of cholinergic neural function after AF64A treatment (see Hanin 1990).

AF64A is used infrequently, largely having been supplanted as a cholinergic neurotoxin by 192-IgG-saporin.

# 5.14 IgG-Saporin

The immunotoxin IgG-saporin consists of (1) a monoclonal antibody directed towards p75<sup>NGF</sup>, the low-affinity membranous receptor for nerve growth factor (NGF), coupled with (2) saporin, a ribosome-inactivating protein derived from seeds of Saponaria officinalis (soapwort) (Wiley 1992; Wiley et al. 1991).

## 5.14.1 Selectivity of IgG-Saporin

IgG-saporin is a selective toxin for cholinergic neurons in basal forebrain, by virtue of their expression of the p75<sup>NGF</sup> receptor on the nerve membrane (Book et al. 1994). Once IgG-saporin is complexed to the p75<sup>NGF</sup> receptor, saporin is internalized by receptor-mediated endocytosis and then transported to the cell body where it enzymatically inactivates ribosomes to inhibit protein synthesis and produce neuronal cell death. IgG-saporin produces near-total destruction of the CBF (see Wiley 1992).

In rat basal forebrain, the cholinergic tract (CBF) is the only one expressing p75<sup>NGF</sup> on the plasma membrane. IgG-saporin thereby selectively destroys these

neurons and thereby produces cholinergic denervation of the CBF target regions – neocortex, hippocampus, amygdale, and olfactory bulb (Rossner et al. 1995; Schliebs 1998; Walsh et al. 1996). Other neuronal tracts of the forebrain survive IgG-saporin treatment because they lack the  $p75^{NGF}$  receptor; this includes striatal cholinergic interneurons and GABAergic neurons within the striatum (Heckers et al. 1994; Wiley et al. 1995).

However, in the cerebellum, the subset of Purkinje cells, which express p75<sup>NGF</sup> receptors during development and into adulthood (Pioro and Cuello 1988), are damaged by IgG-saporin treatment (Berger-Sweeney et al. 1994; Heckers et al. 1994).

## 5.14.2 Effect of IgG-Saporin Lesioning

The CBF is the neuronal tract that degenerates in Alzheimer disease (AD), and the extent of CBF loss correlates with the degree of dementia (Perry et al. 1978). Because IgG-saporin selectively destroys the CBF, this neurotoxin is used to model AD (Schliebs et al. 1996; Wiley et al. 1995).

The neurotoxicity of IgG-saporin has been described in numerous reviews: Bassant et al. 1998; Petrosini et al. 2013 [this *Handbook of Neurotoxicity*]; Rossner 1997; Schliebs et al. 1996; Walsh and Potter 1998; Wiley (1992); Wrenn and Wiley (1998).

## 5.15 3-Nitropropionic Acid (3-NP)

3-Nitropropionic acid (3-NP) ( $O_2NCH_2CH_2COOH$ ), a suicide (irreversible) inhibitor of succinate dehydrogenase (Alston et al. 1977), is a plant and fungal neurotoxin with relative selectivity for basal ganglia (caudate and putamen in humans; striatum in rodents) (Ludolph et al. 1991), although high-dose 3-NP additionally damages hippocampus. In humans who consumed foods with 3-NP and in nonhuman primates treated with 3-NP, the spectrum of neurodegeneration and motor disability, with overall spectrum of effects, resembles/models Huntington's disease (HD).

## 5.15.1 3-NP Mechanism of Action

Coles et al. (1979) identified 3-NP as a suicide inhibitor of succinate dehydrogenase (Huang et al. 2006). Gould and Gustine (1982) first showed that 3-NP produced overt neurotoxicity in basal ganglia.

As succinate dehydrogenase is the major element of mitochondrial respiratory chain complex II, 3-NP inactivation of this site leads to reduced ATP formation, cellular energy depletion, and ultimate preferential degeneration of medium spiny neurons in basal ganglia. NADPH-diaphorase-positive aspiny neurons and cholinergic interneurons survive 3-NP treatment (Beal et al. 1993; Brouillet et al. 1993). Neurodegenerative effects of 3-NP arise when there is prolonged partial (>50%) SDH inhibition (Alexi et al. 1998; Brouillet et al. 1998), resulting in loss of the mitochondrial membrane potential (Leventhal et al. 2000; Maciel et al. 2004), cytochrome c release from mitochondria with calpain activation (Bizat et al. 2003a, b; Nasr et al. 2003), caspase-3 activation (Bizat et al. 2003b),

translocation of Bcl-2, and ultimate neuronal apoptosis (Galas et al. 2004). Calpain activation is considered to be a signature event. Notably, 3-NP-induced calpain activation is preferential to basal ganglia, with calpain remaining active in neocortex and other brain regions where neurodegeneration is generally absent (Galas et al. 2004).

The above processes are also associated with the generation of cellular ROS including  $O_2^{-}$ , HO<sup>•</sup>, and NO formation (Schulz et al. 1995, 1996; Fontaine et al. 2000). Compromise of the energy status of medium spiny neurons is thought to sensitize to glutamate-mediated effects at the NMDA-R, with promotion of glutamate excitotoxicity (Hamilton and Gould 1987; Novelli et al. 1988; Zeevalk and Nicklas 1992; Kim et al. 2000). This latter event was postulated as being responsible for the neurodegeneration with 3-NP and in HD (Albin and Greenamyre 1992; Beal 1992).

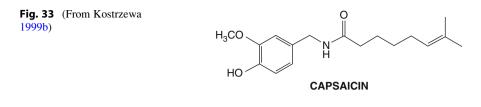
3-NP is commonly administered by osmotic minipumps in rodents, producing dystonia with motor incoordination (Beal et al. 1993; Borlongan et al. 1995). In primates, daily 3-NP administration produces the characteristic striatal neurodegeneration and choreiform movements with cognitive deficits (Brouillet et al. 1995; Palfi et al. 1996). Processes and mechanisms of 3-NP have been described in detail (Brouillet et al. 2005).

#### 5.16 Toxic Vanilloids, TRPV1 Receptor Agonists

#### 5.16.1 Overview

Vanilloids, so-named by virtue of their vanillyl (4-hydroxy-3-methoxybenzyl) moiety, are comprised of three chemical classes: capsaicinoids, resiniferanoids, and diterpenes. Capsaicin (Fig. 33), the best known of the vanilloids, is a capsaicinoid derived from hot chili peppers (Szallasi 1998; Szallasi and Blumberg 1999) – flavoring foods by producing a spicy-hot sensation. The effect is related to activation of TRPV1 receptors (TRPV1-R) on fine-diameter unmyelinated C fibers or thinly myelinated Adelta sensory axons (Holzer 1991) of  $\sim 25$  % of sensory nerves in bipolar dorsal root ganglia and trigeminal ganglia (Liu et al. 1996). Capsaicin, given repeatedly, desensitizes the TRPV1-R (Jancsó and Jancsó 1949; Buck and Burks 1986; Holzer 1991) which undergoes a conformational change in a ligand-gated cation channel (Bevan and Docherty 1993). Additionally, capsaicinoids suppress the expression of peptide neurotransmitters in sensory ganglia (Farkas-Szallasi et al. 1995) while enhancing the expression of galanin, an inhibitor of excitatory neurons (Yanagisawa et al. 1986). High-dose vanilloids are capable of producing overt destruction of sensory neurons (Jancsó et al. 1978).

Insect and animal venoms often contain peptide toxins able to activate the vanilloid receptor (Siemens et al. 2006; Bohlen and Julius 2012) or inhibit it. Capsaicinoids are used in the treatment of neuropathic pain (Rumsfield and West 1991; Winter et al 1995) and are found in "pepper" sprays and in squirrel-free bird food (birds' TRPV1-R lack the vanilloid binding motif).



#### 5.16.2 Mechanism of Vanilloid Action

Vanilloid neurotoxicity is associated with an increase in the Ca<sup>2+</sup> current in sensory neurons, leading to a rise in the cytoplasmic level of Ca<sup>2+</sup> (Chard et al. 1995) and activation of calcineurin and other proteases (Docherty et al. 1996) which dephosphorylate the vanilloid receptor or channel complex and thereby desensitize the receptor complex (Bevan and Docherty 1993). Capsaicin in high concentration (EC<sub>50</sub>, 6.9  $\mu$ M) also reduces the mitochondrial membrane potential (Dedov et al. 2001) and can produce DNA fragmentation, with apoptotic cell death.

The topic of toxic vanilloids is the subject of several reviews (Buck and Burks 1986; Holzer 1991; Bevan and Docherty 1993; Szallasi 1998; Szallasi and Blumberg 1999; Kumar et al. 2013).

## 5.17 Botulinum Neurotoxin

The name botulinum neurotoxin (BoNT) is derived from the Latin "botulus" (i.e., sausage), reflecting discovery of BoNT's ill effects during the Middle Ages – consequent to consumption of poorly prepared or preserved sausage. BoNT and the related tetanus neurotoxin are members of the group of anaerobic bacterial clostridial neurotoxins.

There are seven serological BoNTs, BoNT-A through BoNT-G, with A, B, E, and F subtypes having effects in humans. BoNT consists of a 100kDA heavy chain, paired by a disulfide bond with a 50 kDa light chain. One segment of the BoNT molecule, the binding domain, specifically binds to sites on the choliner-gic nerve, thus imparting selectivity of BoNT neurotoxicity to cholinergic nerves (i.e., motoneurons and parasympathetic nerves). A translocation domain is considered to govern pore formation in the cholinergic plasma membrane, for entry of the catalytic domain into the cholinergic nerve. The catalytic domain inactivates V-snare or T-snare proteins, thus preventing the binding of cytoplasmic vesicles to the cholinergic plasma membrane and thereby preventing exocytosis of the neurotransmitter acetylcholine. BoNT poisoning results in skeletal muscle flaccid paralysis, with death from asphyxiation. However, in regulated dosages, BoNT has proven to be a clinically useful agent for a variety of disorders (Foster et al. 2006; Kostrzewa and Segura-Aguilar 2007; Mehanna and Jankovic 2013; Singh et al. 2013).

Mechanisms, actions, and clinical utility of BoNT are the topic of several recent reviews (Foster et al. 2006; Singh 2006; Kostrzewa and Segura-Aguilar 2007; Mehanna and Jankovic 2013; Singh et al. 2013).

#### 5.18 Quinpirole, a Dopamine D<sub>2</sub>-Receptor Agonist

The definition of a "selective neurotoxin" has become somewhat obscure, when taking the actions of the DA D<sub>2</sub>-receptor agonist, quinpirole, into consideration. Quinpirole, when administered repeatedly – at specified stages of ontogeny – produces long-term sensitization (i.e., a "priming" phenomenon) of the D<sub>2</sub>-R, so that subsequent quinpirole dosing results in exaggerated, sometimes abnormal, behavioral responses. Yet, there is no overt evidence of neurotoxicity, other than the abnormality of behavioral effects to challenge doses of this D<sub>2</sub>-R agonist.

Eilam et al. (1989) first observed that quinpirole challenge in adult rats produced hyperactivity and perseveration in routes (restricted paths in an open environment) without perseveration in movements. Subsequently, ontogenetic postnatal daily quinpirole treatments (2.6 mg/kg/day  $\times$  32 day, base form, ip) of rats were shown to produce enhanced quinpirole-induced behavioral effects. The same abnormal stereotypic response (i.e., licking, grooming, digging, eating) to quinpirole was observed in rats that had been both 6-OHDA lesioned (134 µg, i.c.v., 3 days after birth) as neonates and challenged daily for the first 32 days after birth either with quinpirole or with the DA D<sub>1</sub>-R agonist SKF 38393 (Kostrzewa et al. 1990).

When quinpirole (2.6 mg/kg/day) was administered ip to rats once a day for the first 28 days after birth, quinpirole challenge (25–200 µg) at 8–10 weeks produced a dose-related enhanced yawning response, without there being evidence of a change in the  $B_{max}$  or  $K_d$  for [<sup>3</sup>H]spiperone binding in the striatum (Kostrzewa and Brus 1991). This enhancement of quinpirole-induced yawning was evident at 6 weeks, even in rats that had been primed by low-dose quinpirole (50 µg/kg/day) for 11 consecutive day periods: birth (P0) to P11 (11 days after birth), P12 to P22, or P23 to P33 (Kostrzewa et al. 1993a). In contrast to quinpirole which primes the D<sub>2</sub>-R, the reputedly D<sub>3</sub>-R agonist 7-OH-DPAT [( $\pm$ )-2-(dipropylamine)-7-hydroxy-1,2,3,4-tetrahydronaphthalene] failed to prime a yawning response to either quinpirole or 7-OH-DPAT in rats (Oswiecimska et al. 2000).

Postnatal ontogenetic quinpirole priming (2.6 mg/kg/day) of rats was also associated with adulthood enhancement of a quinpirole (100 or 1000  $\mu$ g/kg)induced antinociceptive effect (hot plate response time) (Kostrzewa et al. 1991). Bizarre quinpirole-induced vertical jumping was observed during the interval of postnatal ontogenetic quinpirole priming (2.6 mg/kg/day, P0–P28) of rats, starting at P18. Quinpirole-induced vertical jumping was evident with quinpirole challenge doses in the range of 0.1–3.0 mg/kg), and the effect of high-dose quinpirole persisted for hours. Notably, vertical jumping was observed only if the cage lid was removed. Otherwise jumping was suppressed, so that rats did not harm themselves by hitting against the cage lid. Typically, paw treading against the cage wall was evident during the duration of the jumping episode (Kostrzewa et al. 1993b). In rats that had been both 6-OHDA lesioned (134  $\mu$ g, i.c.v., 3 days after birth) as neonates and challenged daily from birth with quinpirole, quinpiroleinduced vertical jumping was enhanced (Kostrzewa and Kostrzewa 2012). While changes in striatal D<sub>2</sub>-R number or affinity do not accompany D<sub>2</sub>-R supersensitivity (Kostrzewa and Brus 1991), an acute challenge dose of amphetamine (1.0 mg/kg ip) in 3-month old rats that had been primed with quinpirole during postnatal ontogeny (50  $\mu$ g/kg/day ip, P0–P11) produced a five-fold increase in striatal DA exocytosis, as evidenced in the in vivo microdialysate DA level in awake freely moving rats. In contract, a quinpirole challenge dose in adulthood did not evoke DA exocytosis. In rats that were primed from P0 to P11 by daily quinpirole treatment, there was suppression of striatal DA exocytosis by quinpirole, identical to that observed in control non-primed rats. And in rats that were primed at approximately 2 months of age by 3 increasing doses of quinpirole (25, 50, and 100  $\mu$ g/kg, 1 dose per day), an acute quinpirole challenge dose (100  $\mu$ g/kg) produced even greater suppression of striatal DA exocytosis (Nowak et al. 2001). These findings indicate that D<sub>2</sub>-R supersensitivity has a postsynaptic component (i.e., AMPH-evoked exocytosis).

In neonatally 6-OHDA-lesioned rats (134  $\mu$ g, i.c.v., 3 days after birth) that were additionally primed during postnatal ontogeny with quinpirole (2.6 mg/kg/day, P0–P28), a quinpirole challenge dose (2.6 mg/kg) in adulthood failed to produce an enhanced quinpirole-induced locomotor response. However, quinpirole priming did enhance D<sub>1</sub>-R agonist SKF 38393-induced locomotor responses in adulthood (Brus et al. 2003). This effect is analogous to adulthood heterologous priming of D<sub>1</sub>-R by weekly quinpirole challenge doses in rats that had been lesioned as neonates with 6-OHDA (Criswell et al. 1989).

While DSP-4 neonatal lesioning of noradrenergic innervation prevents ontogenetic quinpirole priming of  $D_2$ -R (Nowak et al. 2006), neonatal co-lesioning with DSP-4 and 5,7-DHT results in enhanced quinpirole-induced and 7-OHDA-induced yawning as well as enhanced apomorphine-induced stereotypies (Nowak et al. 2009). These findings indicate that both noradrenergic and serotoninergic nerves have a prominent influence on  $D_2$ -R priming.

In quinpirole-primed rats, spatial memory deficits and enhanced skilled reaching were observed in adulthood (Brown et al. 2002). Enhanced quinpirole-induced locomotor and stereotyped responses observed in adult rats that were quinpirole primed during postnatal ontogeny are attenuated by acute nicotine pretreatment in adulthood; and this effect was accompanied by increased  $[^{125}I]\alpha$ -bungarotoxin binding in striatum and hippocampus and greater  $[^{3}H]$ cytisine binding in midbrain and cerebellum. These findings indicate that nicotinic  $\alpha_7$  and  $\alpha_4\beta_2$  receptor parameters, respectively, are regionally altered in quinpirole-primed rats (Tizabi et al. 1999). In separate studies, nicotine adulthood treatments (0.3 mg/kg x 2/day) for 14 consecutive days reversed performance deficits in quinpirole-primed rats on the Morris water task and skilled reaching task and partially reversed a 36 % decrease of choline acetylcholine transferase in hippocampus. This finding supports the role of nicotinic receptors in quinpirole priming and projects involvement of cholinergic systems in the priming phenomenon (Brown et al. 2004).

 $D_2$ -R supersensitivity is implicated in obsessive-compulsive disorder (Eilam et al. 1989, 2006; Szechtman and Woody 2004, 2006; Alkhatib et al. 2013) and is

regarded as a major component of schizophrenia in which there is an elevation in the number of  $D_2^{high}$  receptors (Seeman et al. 2005, 2007; Seeman 2011). So-called breakthrough DA-R supersensitivity is also regarded as a factor in antipsychotic treatment failure (Samaha et al. 2007).

The process of D<sub>2</sub>-R supersensitivity, as relating to quinpirole-induced ontogenetic priming, appears to be a lifelong permanent effect, still evident in rats approaching 2 years of age (Brus et al. 1998; Oswiecimska et al. 2000). Quinpirole induction of D<sub>2</sub>-R supersensitivity is discussed in several reviews (Kostrzewa 1995; Kostrzewa et al. 2004, 2008a, 2011; Nowak et al. 2004; Brown et al. 2012). Permanent sensitization of D<sub>2</sub>-R and resulting alteration of behaviors, despite lack of evidence of neuropathology, does nonetheless represent a neurotoxicity.

#### 5.19 Other Neurotoxins

The overview of selective neurotoxin in this chapter is intended to highlight the most commonly known neurotoxins and most commonly used neurotoxins. Not all neurotoxins have been listed and/or described in detail in this overview. There are a number of other EAAs and other types of neurotoxins that are used, and some are described in greater detail in other chapters of the *Handbook of Neurotoxicity*.

## 6 The Future of Neurotoxins in Biomedicine

For the past 50 years, selective neurotoxins have been instrumental in:

- Mapping nerve tracts in the CNS: Destruction of the nerve terminal field is often followed by axonal swelling, retrograde throughout its entire length, back to the perikaryon. A marker for this phenotypic nerve provides an excellent means to map the route of the axons. For example, by lesioning the terminal field for noradrenergic nerves and treating with a fluorescent-tagged antibody for dopamine-beta-hydroxylase, noradrenergic axons will glow brightly green and thereby show the exact course of the noradrenergic nerve tract through the brain.
- 2. Demonstrating nerve fiber sprouting: When neurotoxins destroy one axon of a bipolar (i.e., 2 axons) or multipolar nerve, the more proximal branch(es) of the nerve often sprout and hyperinnervate regions that are normally innervated by the surviving axonal branch(es). At times, the surviving branch(es) extends their geographic domain to innervate new regions which are not normally innervated by this nerve (neo-innervation). Neurotoxins have provided the means to uncover these phenomena. Also, by virtue of selective neurotoxicity, these findings are "pure" not encumbered by complications ensuing, were more than one neuronal phenotype damaged.
- 3. Determining the physiological role played by specific neuronal tracts: As indicated by WB Canon, the function of a nerve can be deduced by denoting altered effect when the nerve is destroyed (Cannon and Rosenblueth 1937). Accordingly, the importance of dopaminergic innervation in nucleus accumbens

with the euphoric and other effects of drugs of abuse can be deduced. The same approach is successfully used to link other nerve phenotypes with a particular brain region and associated behavioral effect.

- 4. Ascribing neuronal denervation with receptor supersensitization phenomena: The proliferation in numbers of receptors following selective destruction of a nerve can be reasonably quantified; and possible changes in the affinity of these receptors can be studied. Also, the relative contribution of high- and low-affinity states of the receptor can be elucidated. Selective neurotoxins are especially valuable for such studies. However, since the time that selective neurotoxins have been employed for such studies, it has become clear that receptor supersensitization (i.e., ability of the receptor to produce an exaggerated response) is not necessarily related to a change in receptor number. Additional receptors do not necessarily correlate with increased receptor sensitization.
- 5. Characterizing specific neuronal phenotypes with human neurological and psychiatric disorders: Selective lesioning of particular nerve types or specified nerve tracts has been of immense value in animal modeling of human neuropathological and psychiatric states. Ideally, one seeks (a) face validity (mirroring of a behavioral deficit), (b) construct validity (conforming with a pathophysiological basis of a disorder), and (c) predictive validity (ability to project a therapeutic intervention) in an animal model. Some models relate to merely "predictive validity." For example, perinatal 6-OHDA treatment of rats leads to an adolescent and adulthood hyperactivity and attention deficit, which is responsive to amphetamine just as human ADHD is responsive to amphetamine (see Kostrzewa et al. 2008b). Yet, human ADHD has no relationship with near-total dopaminergic denervation of forebrain as found in this animal model. Other animal modeling of human disorders is reliant on use of selective neurotoxins.
- 6. Identifying intraneuronal mechanisms and processes that lead to cell death by apoptosis, necrosis, and necroptosis: Neurotoxins commonly evoke a number of processes that lead to cellular instability and compromised viability. Classically there is an increase in intracellular Ca<sup>2+</sup> levels, also in the number of ROS and reactive nitrogen species, changes in oxidative phosphorylation accompanied by altered mitochondrial membrane potential, mitochondrial release of cytochrome c into the cytoplasm, increased phosphorylation of proteins, promotion of aggregates of abnormal proteins, activation of Bax and other proapoptotic elements along with suppression of bcl and other protective cellular elements, translocation of proteins to the nucleus, and other changes. Neurotoxins have aided immensely in the study of cell death phenomena.
- 7. Recognition of the role played by microglia, astrocytes, and immune cells in neurodegeneration: Selective neurotoxins have been instrumental in evoking neuronal damage and thereby provoking nonneural cellular responses to injury. These include astrocytes, activated microglia, and immune cells which arise in a uniquely timed and ordered array. The incentive to understand mechanisms of neurotoxicity has aroused interest in the role played by other cells in neuropathological processes.

8. Broadening the understanding that there are multiple cellular targets whereby drugs may intervene to disrupt an otherwise programmed cell death: There was an awakening of interest in new means by which drugs can conceivably target one or another process to facilitate recovery of an injured neuron. In addition, there is now global interest in neuroprotectants, which can prevent an otherwise deleterious effect on nerves.

Looking ahead we can foresee the introduction of multiple kinds of selective neurotoxins, in order to broaden the horizon for these agents in elucidating the association between specified nerve and nerve tracts with biological and behavioral actions. We can envision:

- 1. An increase in the number of phenotypically selective neurotoxins
- 2. Introduction of neurotoxins that act on novel intracellular targets
- 3. More neurotoxins tagged to antibodies
- 4. Trojan horse botulinum toxins, engineered with binding proteins that target an ever-expanding field of susceptible cells, and conveying a toxic load into these cells
- 5. Engineered immunotoxins, altered to target specified cells analogous to IgG-saporin
- Agonists/antagonists directed towards neurotrophin receptors, so as to compromise development of specified neuronal phenotypes
- Antibodies for specific neurotrophins, analogous to that of anti-NGF, producing immuno-disarray of specified types of nerves
- 8. Alkylating agents analogous to DSP-4
- 9. Added excitatory amino acids (EAAs)
- 10. Inhibitors of (endogenous) neuroprotectants
- 11. Agents that desensitize nerves, akin to capsaicin
- 12. A host of agents that act by unanticipated means

In the process, there will be a discovery of novel neuroprotectants, some with targeted specificity and others with generalized effects. As more becomes known of cellular processes and mechanisms in the neuronal response to injury, so too does the number of possible neuroprotectants increase.

We cannot foresee changes that will occur in the next century. So many more revolutions are certain, and someone looking back 100 years from now will likely be confused by what will probably seem to be limited scope in the assumed "direction" of this field. When tomorrow sees today's views as too narrow in scope, this will be proof that we, today, recognized how important the topic of neurotoxins is, for the development of neuroscience and an ever increasing understanding of how nerves work (adapted from Kostrzewa 1999a).

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# Necroptosis, a Potential Therapeutic Target for Neurological Disorders

# Jing Chen, Richard M. Kostrzewa, and Xingshun Xu

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J. Chen • X. Xu (🖂)

R.M. Kostrzewa

Department of Biomedical Sciences, Quillen College of Medicine, East Tennessee State University, Johnson City, TN, USA e-mail: KOSTRZEW@mail.etsu.edu

Department of Neurology, The Second Affiliated Hospital of Soochow University, Suzhou City, Jiangsu, China

The Institute of Neuroscience, Soochow University, Suzhou City, Jiangsu, China e-mail: jing\_ch.china@hotmail.com; Xingshunxu@suda.edu.cn

#### Abstract

Necrosis is considered to be an unregulated and chaotic cell death. However, recent advances in cell death strategies support necroptosis as a form of regulated programmed necrotic cell death. In response to TNF- $\alpha$  or Fas ligands, necroptosis can be induced by cell death receptors in multiple cell lines in the presence of a caspase inhibitor z-VAD; necroptotic cell death has been found to play an important role in normal development, immunity, inflammation, cancer, and human diseases. In this chapter, the molecular mechanisms governing necroptosis, recent findings about the upstream and downstream schema of necroptosis, and potential therapeutic targets in neurological disorders are discussed. After being activated by TNF- $\alpha$  (or Fas ligands) and death receptors, receptor-interacting proteins 1 and 3 (RIP1 and RIP3) form a complex, which play a central role in the induction of necroptosis. RIP3 phosphorylates and activates mitochondrial proteins mixed lineage kinase domain-like protein (MLKL) and PGAM5, resulting in the execution of necroptosis by dynamin-related protein 1, the GTPase that controls mitochondrial fission. Some small molecules such as necrostain-1 and necrosulfonamide target different steps of necroptosis and impede the progress of necroptosis. FADD, caspase-8, CLIP, and CYLD positively or negatively regulate RIP1-/RIP3-dependent necroptosis by different mechanisms. Recent studies demonstrate the involvement of necroptosis in many neurological disorders including stroke, trauma, neonatal hypoxic-ischemic encephalopathy, and Huntington's disease. As a potential therapeutic target, the understanding of necroptotic mechanisms will provide new insights to develop more potent neuroprotectants and specific therapeutic strategies for clinical treatments of neurological disorders.

Keywords			
Necroptosi	is • Neurological diseases • RIP1 • RIP3		
List of Abbreviations			
AIF	Apoptosis-inducing factor		
ANT	Adenine nucleotide translocase		
Aur-A	Aurora-A kinase		
BHA	Butylated hydroxyanisole		
CCI	Controlled cortical impact		
cIAP1/2	Apoptosis protein 1/2		
СурА	Cyclophilin A		
CypD	Cyclophilin D		
DRP1	Dynamin-related protein 1		
FADD	Fas-associated protein with death domain		
FLIPL	FLICE-like inhibitory protein long		
GLUD1	Glutamate dehydrogenase 1		
GLUL	Glutamate–ammonia ligase		
HI	Hypoxia–ischemia		

HIE HMGB1 HSP90 IAPs MEFs MKRN1 MLKL OGD Plk1 RHIM RIP ROS TNEP1	Hypoxic-ischemic encephalopathy High-mobility group box 1 protein Heat shock protein Inhibitors of apoptosis Mouse embryonic fibroblasts Makorin Ring Finger Protein 1 Mixed lineage kinase domain-like protein Oxygen-glucose deprivation Polo-like kinase 1 RIP homotypic interaction motif Receptor-interacting protein Reactive oxygen species TNE recentor 1
ROS	
TNFR1	TNF receptor 1
TNF-α	Tumor necrosis factor- $\alpha$
z-VAD	z-VAD-fmk

### 1 Introduction

Neurological disorders, from the gamut of acute medical emergencies such as stroke to debilitating chronic neurodegenerative disorders such as Parkinson's disease and Alzheimer's disease, are related to neuronal cell damage, cell repair, and final cell death or survival. For the prevention or therapeutic intervention for these neurological disorders, the process of cell death still is one of major targets.

Neuronal loss is observed in many neurological disorders including stroke, Parkinson's disease, and Alzheimer's disease (Ekshyyan and Aw 2004). However, the kinds of cell death existing in the pathological process of the multiplicity of neurological diseases are still not clear. On the basis of intracellular morphological changes, it was considered that there were three types of cell death in embryonic developmental processes (Schweichel and Merker 1973). Type I cell death is characterized by cytoplasmic shrinkage, pyknosis, nuclear condensation, and late cell fragmentation, which is identical to "apoptosis," as described by Kerr (Kerr et al. 1972). Type II cell death is characterized by substantial autophagic vacuolization in the cytoplasm, termed lately as "autophagy." Type III cell death is characterized by clumping of chromatin, swelling of organelles, and cell disintegration including the rupture of nuclei, mitochondria, lysosomes, and plasma membrane; and this is considered to be "necrosis" (Kitanaka and Kuchino 1999). In the past decades, accumulating evidences demonstrate that many alternative cell death mechanisms exist during the development and pathology of diseases including necroptosis, elimination by shedding, keratinocyte death by cornification, and cell-cell cannibalism by entosis (Yuan and Kroemer 2010). Among the reported forms of cell death, "apoptosis," "autophagy," and "necroptosis" are regarded as the major types most closely associated with neuropathology in neurological and neurodegenerative disorders

(Edinger and Thompson 2004). Apoptotic and autophagic cell death regulation pathways have been extensively studied and well established in the past two decades. Unlike apoptosis and autophagy – the forms of programmed cell death – necrosis has long been considered as an uncontrolled nonprogrammed cell death characterized by rapid chaotic breakdown of the cell under intolerable conditions. More recently, necrosis has been recognized as a programmed and regulated cell death process (Degterev et al. 2005; Edinger and Thompson 2004). Since necroptosis – a form of necrotic cell death – was proposed in 2005 (Degterev et al. 2005), the concept of regulated necrosis stirred cell death research, while regulation of signal pathway transduction became one of the hot topics in biomedical research.

Necroptosis represents a specific form of caspase-3-independent programmed necrotic cell death that is induced by cell death ligands such as TNF- $\alpha$  and Fas ligand via cell death receptors in multiple cell types, in the presence of an apoptosis inhibitor (Degterev et al. 2005). The increasing evidences have demonstrated the existence of necroptosis in a variety of neurological disorders. In this chapter, the regulative pathway of necroptosis and its potential therapeutic application in neurological disease will be discussed.

# 2 The Cellular Signal Transduction of Necroptosis

Necroptosis, as a form of necrotic cell death, was identified by Degterev and his colleagues based on the observation of cell death induced by TNF- $\alpha$  or Fas ligand in the presence of an apoptosis inhibitor (Degterev et al. 2005). Initially, necroptosis was not considered to share signal pathways with apoptosis and was thereby accorded as an independent cell death process distinct from apoptosis. However, recent advances have shed greater insight into the mechanisms by which necroptosis is regulated and executed. Actually, as an alternative form of cell death, necroptosis can not only cross talk with apoptosis and autophagy but under some circumstances can also recruit other cell death processes such as apoptosis or autophagy (Duprez et al. 2012; Lu and Walsh 2012; Simenc and Lipnik-Stangelj 2012; Vandenabeele et al. 2010; Zhang et al. 2009). In most pathological situations, different kinds of cell death may exist and lead to both cell loss and tissue damage (Long and Ryan 2012; Yahiro et al. 2012). However, one form of cell death will dominate the pathological process. When the signal pathways of this dominant cell death process are blocked (e.g., caspase inhibitor block of apoptosis), an alternative cell death process such as necroptosis can be increased and become the dominant form of cell death (Han et al. 2011). By understanding the conversion or switch between different forms of cell death, it becomes possible to identify new clinical targets - thus enabling the advance of new therapeutic strategies. As for necroptotic cell death, the deciding factors for the conversion from/to apoptosis are the central protein complexes RIP1 and RIP3; but positive regulators (such as deubiquitinase CYLD) and negative regulators (such as caspase-8, FADD) also closely control the fate of cells (survival, necroptosis, or other forms of cell death converted from necroptosis). The advances on regulation of necroptosis are discussed below.

#### 2.1 The Central Signals for Necroptosis: RIP1 and RIP3

Receptor-interacting protein (RIP) kinase family has been identified as having seven members so far (Zhang et al. 2010). RIP kinase family members share a homologous kinase domain but have different functional domains. Since their discovery as a novel form apoptosis-inducing protein, RIP kinases have emerged as essential sensors of cellular stress (Stanger et al. 1995). RIP kinases not only trigger pro-survival through the activation of transcription factors such as NF- $\kappa$ B and AP-1 but also initiate the death-inducing programs (Meylan and Tschopp 2005). Hence, RIP kinases are regarded as crucial regulators of cell survival and death (Festjens et al. 2007). Among these family members, RIP1 and RIP3 are specifically related to necroptosis (Cho et al. 2009) (Fig. 1).

RIP1, the first member of the RIP kinase family, contains three domains: an N-terminal kinase domain, an intermediate domain, and a C-terminal death domain. Although the death domain of RIP1 was shown to bind to several death receptors, such as TNFR1 to activate NF- $\kappa$ B and induce apoptosis (Holler et al. 2000; Stanger et al. 1995), the serine/threonine activity of RIP1 is essential for necroptosis (Degterev et al. 2005). Moreover, RIP1 is regarded as a crucial adaptor kinase and a switch between different forms of cell death in response to extracellular stress or damaging stimuli such as TNF- $\alpha$  and Fas ligands (Festjens et al. 2007). TNF- $\alpha$  is the most extensively studied extracellular signal that leads to apoptosis or necroptosis. Under the apoptosis situation, TNF- $\alpha$  stimulation sequentially induces the formation of two protein complexes: complex I and complex II. In the cytoplasmic membrane, TNF- $\alpha$  binds to TNF receptor 1 (TNFR1) and induces the formation of protein complex I including TRADD, TRAF2, RIP1, and cIAP1. RIP1 is recruited to the protein complex via its death domain to TNFR1. After the formation of protein complex I, these proteins dissociate from TNFR1, and RIP1 enters the cytosol. In the cytosol, RIP1 forms protein complex II with caspase-8, Fas-Associated protein with Death Domain (FADD), and other proteins to activate the caspase cascade to execute apoptosis. In the presence of pancaspase inhibitor z-VAD-fmk (z-VAD), caspase-8 activity is inhibited and the apoptotic pathway is blocked. In this situation, RIP1 interacts with RIP3 to form Ripsome and initiate necroptosis as an alternative cell death. In the apoptosis-deficient condition, the expression of an inactive RIP1 or the inhibition of RIP1 activity by Nec-1 inhibits TNF-induced necroptosis (Degterev et al. 2005; Holler et al. 2000). Mice lacking RIP1 appear normal at birth but fail to thrive, displaying extensive apoptosis in both the lymphoid and adipose tissues and dying at 1–3 days of age (Kelliher et al. 1998). Therefore, RIP1 kinase activity is crucial for this alternative cell death pathway of necroptosis.

The increasing body of evidences demonstrate that RIP3 is a determinant for necroptosis (Challa and Chan 2010). Recent studies show that cells which lack RIP3 expression are resistant to necroptosis, despite the normal RIP1 expression (Cho et al. 2009; He et al. 2009). When RIP3 expression was restored with an shRNA-resistant RIP3 transgene, necroptotic cell death was reestablished (He et al. 2009). However, if RIP3 Kinase-Dead Mutant K50A abolishes RIP3 kinase

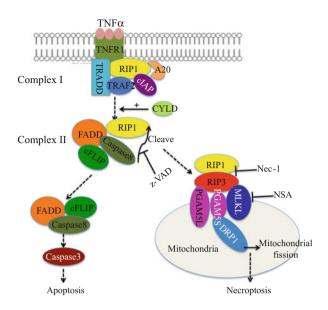


Fig. 1 TNF- $\alpha$ -elicited signaling pathways on apoptosis and necroptosis. Upon TNF- $\alpha$  stimulation, TNFR1 induces the formation of protein complex I including TRADD, TRAF2, and RIP1 at the plasma membrane. RIP1 is modified by polyubiquitination via cIAP1 and A20. After the dissociation from TNFR1, RIP1 enters the cytosol and deubiquitinates by CYLD. In the cytosol, RIP1 forms protein complex II with FADD, caspase-8, and cellular FLIP (cFLIP). The composition of protein complex II and its activation determine the directions of the cell death: apoptosis or necroptosis. FADD contains a death-effector domain, which recruits and causes the autocatalytic activation of the initiator caspase-8; caspase-8 directly cleaves intracellular substrates or activates other caspases such as caspase-3, finally resulting in apoptosis. The apoptotic machinery FADD, caspase-8, and cFLIP suppress necroptosis via the cleavage of RIP1 by activated caspase-8. In apoptosis-deficient conditions such as in the presence of pancaspase inhibitor z-VAD, caspase-8 activation is inhibited and blocks the RIP1 cleavage. RIP1 then interacts with RIP3 to form a RIP1-RIP3 complex. RIP3 phosphorylates two proteins in the outer mitochondrial membrane: a protein kinase MLKL and a phosphatase PGAM5. After the activation of MLKL, short form of PGAM5 (PGAM5S) activates DRP1, the GTPase that controls mitochondrial fission, in a hydrophobic environment, leading to mitochondrial fission and necroptosis. The necroptosis inhibitors necrostain-1 (Nec-1) and necrosulfonamide (NSA) block necroptotic cell death by targeting the interaction of RIP1 and RIP3 and MLKL, respectively

activity, necroptosis remains blocked, indicating that the kinase domain is still essential for the role of RIP3 (He et al. 2009).

The interaction of RIP1 and RIP3 is required for mutual and autophosphorylation of RIP1 and RIP3 and the activation of the downstream necrotic signaling pathway (Cho et al. 2009; He et al. 2009). Both the intermediate domain of RIP1 and the unique C-terminus of RIP3 have a RIP homotypic interaction motif (RHIM), which is required for mediating the interaction between RIP1 and RIP3 (Meylan and Tschopp 2005). The small-molecule necrostain-1, an inhibitor of RIP1 kinase activity, blocks necroptosis by impairing the interaction between RIP1 and RIP3 (Cho et al. 2009; He et al. 2009). The mutation in the

RHIM domain of RIP proteins inhibits necroptotic cell death (Cho et al. 2009), suggesting that the interaction of RIP1 and RIP3 or RIP1–RIP3 complex is crucial for death receptor-mediated necroptosis. The apoptotic pathway seems to be controlled by a caspase network, whereas necroptotic signaling is controlled by a kinase cascade and highly regulated by an intracellular protein platform (Gunther et al. 2013).

# 2.2 The Upstream Positive and Negative Regulators of Necroptosis

#### 2.2.1 FADD/Caspases-8/FLIP

FADD is an adaptor molecule that bridges the Fas receptor or other death receptors such as TNFR1 to caspase-8 through its death domain to form the death-inducing signaling complex during apoptosis (Bodmer et al. 2000; Kreuz et al. 2004). FADD containing a death-effector domain can recruit and cause the autocatalytic activation of the initiators caspase-8 and caspase-10, which can directly cleave intracellular substrates or activate other caspases including caspase-3 (Tourneur and Chiocchia 2010). When caspase-8 is activated, it mediates apoptosis; at the same time, caspase-8 prevents necroptotic cell death by cleaving RIP1 and/or RIP3 or by binding FLICE-like inhibitory protein long (FLIPL) to form an "inhibitory" caspase-8-FLIPL heterodimer (Lu et al. 2011; Oberst et al. 2011; Rebe et al. 2007). While caspase-8 is inactivated by a pancaspase inhibitor z-VAD or caspase-8 knockout, the proteolytic inactivation of RIP1 and RIP3 is prevented and the inhibitory effect of caspase-8 on necroptosis is abolished. These are confirmed by in vitro and in vivo experiments. In antigen-activated T cells, the ablation of caspase-8 or FADD can lead to RIP1-dependent necroptosis (Bell et al. 2008; Ch'en et al. 2008). Similarly, caspase-8- and FADD-deficient mice demonstrated severe embryonic phenotype including cardiac malformations and hemorrhage mediated by RIP1- and RIP3-dependent necroptosis (Gunther et al. 2013; Kaiser et al. 2011). Interestedly, caspase-8/RIP1 or caspase-8/RIP3 double knockout mice are viable and survive into fertile adults; RIP1 or RIP3 knockout rescues embryonic lethality phenotype observed in caspase-8- or FADD-deficient mice (Kaiser et al. 2011; Oberst et al. 2011; Zhang et al. 2011a). These experiments demonstrate the regulation of caspase-8 and FADD on RIP1- and/or RIP3-mediated necroptosis.

FLIP has a similar structure to caspase-8, but lacks the catalytic activity. FLIP may be involved in the regulation of necroptotic cell death via RIP kinase, as FLIP knockout mice have a similar phenotype to that of caspase-8- and FADD-deficient mice (Yeh et al. 2000). However, in a recent study, RIP3 ablation rescued embryonic phenotype of FADD-deficient mice, but did not rescue the phenotype of FLIP-deficient mice (Dillon et al. 2012), suggesting distinct functions of FLIP from FADD and caspase-8 on the regulation of necroptosis. Surprisingly, embryos at E9.5–E10 in the FLIP and RIP3 double knockout mice display obvious apoptotic cell death (Dillon et al. 2012), demonstrating the antiapoptotic activity of FLIP on FADD-/caspase-8-mediated apoptosis. Therefore, in the absence of FLIP, caspase-8

may form homodimers to activate apoptotic pathways and result in cellular apoptosis; in the presence of FLIP, the heterodimer of caspase-8 and FLIP forms a complex with FADD to prevent RIP kinase-dependent necroptosis (Dillon et al. 2012; Feoktistova et al. 2011). There is constant interplay between death and survival complexes, which compete with each other until one eventually dominates and determines the cell's fate (Declercq et al. 2009). The regulation of FADD/caspase-8/FLIP on necroptosis and apoptosis demonstrates that various death pathways are closely intertwined, and cellular processes are in tight control of each other.

# 2.2.2 Deubiquitinase (CYLD/A20/Cezanne) and Ubiquitinase (IAPs/MKRN1)

Protein ubiquitylation and deubiquitylation are important processes to regulate numerous signal pathways and proteins. Ubiquitylation status of RIP1 determines the type of RIP1-mediated cell death. When RIP1 is recruited to TNFR1, RIP1 is heavily modified by ubiquitination; however, ubiquitinated RIP1 was not found in the RIP1–RIP3 complex (Cho et al. 2009; Teng et al. 2005). The different ubiquitination patterns imply that RIP1 is deubiquitinated in the process of necroptosis. Therefore, the enzymes for ubiquitination and deubiquitination of RIP1 protein are the regulators of necroptosis.

CYLD is a lysine-63 (K63)-specific deubiquitinase, which was identified as an important regulator of necroptosis in a genome-wide RNAi screen (Hitomi et al. 2008). Knockdown of CYLD inhibits TNF- $\alpha$ -induced necroptosis in cultured human Jurkat cells (Hitomi et al. 2008), which suggests that CYLD may be responsible for the deubiquitination of RIP1 during necroptosis. A recent study confirmed that RIP1 is a substrate of CYLD (Wright et al. 2007). Consistent with this, CYLD promotes assembly of the RIP1-FADD-caspase-8 complex in response to TNF- $\alpha$  treatment, indicating that deubiquitination of RIP1 by CYLD is an essential step in this process (Wang et al. 2008). CYLD was recently found to be cleaved by caspase-8 and thereby prevents necroptosis in mouse embryonic fibroblasts (MEFs) upon TNF-a stimulation; however, overexpression of a noncleavable CYLD mutant rapidly promotes necroptotic cell death in response to TNF- $\alpha$  treatment (O'Donnell et al. 2011), which connects CYLD, caspase-8, and necroptosis together. Therefore, RIP1 polyubiquitylation may hinder the recruitment of downstream proteins of necroptosis such as RIP3; CYLD facilitates necroptosis by removal of the polyubiquitin chains on RIP1. The positive regulation of CYLD on necroptosis is also controlled by caspase-8 activity, a negative regulator of necroptosis. Similar to CYLD, ubiquitin-editing enzymes A20 and Cezanne are also reported to deubiquitinate RIP1 protein (Enesa et al. 2008; Shembade et al. 2010).

#### 2.2.3 cIAPs

Inhibitors of apoptosis (IAPs) characterized by a variable number of highly conserved baculoviral IAP repeat motifs including cellular inhibitor of apoptosis protein 1/2 (cIAP1/2) and X-linked inhibitor of apoptosis protein (XIAP) have been thought to function primarily to inhibit apoptosis. A "Really Interesting New Gene" (RING) domain with ubiquitin protein ligase (E3) activity in the C-terminal of IAPs implies that IAPs are multifunctional proteins to regulate the modification of its target protein via ubiquitylation. Recent studies demonstrate that IAPs also regulate RIP1 ubiquitination and RIP1-dependent necroptotic cell death (Bertrand et al. 2008; McComb et al. 2012; Tenev et al. 2011). IAPs can directly conjugate polyubiquitin chains to RIP1 and block RIP1-dependent ripoptosome formation containing the core components RIP1, FADD, and caspase-8 by targeting the components of ripoptosome for ubiquitylation and degradation. Deletion of IAPs enhances ripoptosome formation and the association of RIP1 and caspase-8, resulting in ripoptosome-mediated apoptosis or necroptosis (Tenev et al. 2011). In tumor cells, cIAP1 inhibition by its antagonists or RNA interference facilitates TNF-induced necroptosis (He et al. 2009; Vanlangenakker et al. 2011). Consistent with this, McComb et al. found that cIAPs also regulate RIP3 via ubiquitination to inhibit the RIP1-RIP3 death complex, and reduced cIAP activity in vivo by the exposure to SMAC mimetics or cIAP1/2 knockout leads to elevated macrophage necroptotic cell death (McComb et al. 2012). Therefore, cIAPs are important targets for preventing apoptosis and necroptosis in acute and chronic diseases or inducing cell death in cancer therapy.

# 2.2.4 Ubiquitinase on FADD-Makorin Ring Finger Protein 1 (MKRN1) E3 Ligase

A recent study shows that FADD regulates cell death and survival via a ubiquitination-induced degradation pathway (Lee et al. 2012). Makorin Ring Finger Protein 1 (MKRN1) E3 ligase, a ubiquitinase, ubiquitinates FADD; ubiquitinated FADD is subjected to degradation through a proteasomal pathway. The knockdown of MKRN1 increases the stability of FADD and facilitates caspase-8- and caspase-3-mediated apoptosis in response to death signals; however, the overexpression of MKRN1 suppresses extrinsic apoptosis by increasing FADD degradation (Lee et al. 2012). In the presence of caspase inhibitors, MKRN1 knockdown significantly increases RIP1–RIP3–FADD complex formation and enhances necroptotic cell death upon TNF- $\alpha$  and z-VAD treatment (Lee et al. 2012). This suggests that ubiquitinase MKRN1 also regulates apoptosis and necroptosis via the ubiquitination and degradation of FADD.

#### 2.2.5 Polo-Like Kinase 1 (Plk1) and Aurora-A Kinase (Aur-A)

Plk1 and Aur-A are two important mitotic kinases in the regulation of multiple mitotic processes (Macurek et al. 2009). A recent study has shown that Plk1 and Aur-A are involved in the phosphorylative modification of FADD and cooperatively phosphorylate FADD at S194 and S203, respectively (Jang et al. 2011). The double-phosphorylated FADD results in its dissociation from RIP1, which is activated to necroptotic cell death; however, the single-phosphorylated FADD cannot induce cell death, suggesting that double phosphorylation of FADD is critical for RIP1-dependent necroptosis (Jang et al. 2011). Therefore, inhibition of double phosphorylation of FADD by mitotic kinases Plk1 and Aur-A may block necroptosis. In contrast, another study indicates that the small-molecule BI2536, an

inhibitor of Plk1, increases PARP-1-mediated necroptosis (Deeraksa et al. 2013). This inconsistency of two studies may be explained by the specificity of small molecules on Plk1 and the difference of cell lines. However, both studies demonstrate the involvement of mitotic kinase Plk1 on the regulation of necroptosis. More information on how Plk1 regulates RIP1-dependent necroptosis should be investigated in future studies.

#### 2.3 Downstream Signals of Necroptosis

While the upstream signaling of RIP1–RIP3 and RIP1 ubiquitination are important to determine how necroptosis is initiated, the identification of downstream components is necessary to determine how necroptosis is executed. Caspases are the executioners for apoptotic cell death; however, these proteases have no positive role in necroptosis (Pop et al. 2011). Thus far, an analogous class of executioner proteins has not been identified for necroptosis. Some downstream signals relevant to necroptosis are reported, and this information may provide clues for future studies to unveil necroptosis.

#### 2.3.1 Mixed Lineage Kinase Domain-Like Protein (MLKL)

MLKL belongs to the protein kinase superfamily, since a kinase domain is identified within the C-terminus of the protein; however, its function and substrates are still unknown. Recently, two research groups independently demonstrated that MLKL is a substrate of RIP3 and the direct downstream signal of RIP1-RIP3 complex-mediated necroptosis (Sun et al. 2012; Zhao et al. 2012). Sun and colleagues identified MLKL as an interacting protein of RIP3, by immunoprecipitation and mass spectrometry analysis (Sun et al. 2012). Knockdown of MLKL prevents the enlarged RIP3 punctae during necrosis progress and arrests necrosis at a specific step, which is an effect identical to that caused by necrosulfonamide, a newly identified necroptosis inhibitor and MLKL inhibitor (Sun et al. 2012). Their further findings show that the interaction of MLKL and RIP3 is initiated by autophosphorylation of RIP3, and then two critical residues of MLKL for necrosis are phosphorylated at threonine 357 and serine 358 by RIP3 (Sun et al. 2012). Zhao et al. also demonstrated that MLKL is a key downstream mediator of necrosis directly regulated by RIP3 kinase (Zhao et al. 2012). By a kinase/phosphatase shRNA library screening in a human colon adenocarcinoma cell line, they found that a lack of MLKL blocks TNF-induced necrosis but does not affect the interaction of RIP1 and RIP3, suggesting that MLKL is the downstream of RIP1-RIP3 complex (Zhao et al. 2012). Further findings indicate that MLKL is recruited to RIP1–RIP3 complex by interacting with RIP3, but not mutant RIP3 without kinase activity (Zhao et al. 2012), which is consistent with Sun et al.'s results (Sun et al. 2012). Although MLKL is proven to be directly downstream of RIP3, the downstream of signals of MLKL remains elusive.

#### 2.3.2 PGAM5

PGAM5 is a mitochondrial protein phosphatase in the outer membrane of the mitochondria. A recent study found that PGAM5 coprecipitates with RIP3 under necroptosis-inducing conditions (Wang et al. 2012). More importantly, RIP3 phosphorylates PGAM5 and consequently increases PGAM5 phosphatase activity (Wang et al. 2012), suggesting that PGAM5 is also the substrate of RIP3. Knockdown of either splice variant of PGAM5 (PGAM5L or PGAM5S) by siRNA reduces necroptosis induced by different stimuli, indicating that both isoforms of PGAM5 are essential for necroptosis; however, only PGAM5S phosphorylation is dependent on the presence of MLKL and is blocked by the MLKL inhibitor necrosulfonamide (Wang et al. 2012). These findings suggest that PGAM5S is also recruited to the RIP1-RIP3-MLKL complex. Further study indicates that PGAM5S is phosphorylated after the necrosome (RIP1, RIP3, MLKL, and PGAM5L) is shifted to a more hydrophobic environment that contains dynaminrelated protein 1 (DRP1), the GTPase that controls mitochondrial fission. Interestingly, knockdown of DRP1 prevents necroptosis, and knockdown of PGAM5S attenuates DRP1 activation (Wang et al. 2012), demonstrating that DRP1 is required for necroptosis and is regulated by PGAM5S. Therefore, PGAM5S is also a substrate and downstream signal of RIP1-RIP3 complex, and DRP1 may be one of the executioners in the necroptotic pathways.

# 2.3.3 Poly(ADP-Ribose) Polymerase-1/Apoptosis-Inducing Factor (PARP-1/AIF) Pathway

PARP is a family of nuclear enzymes that catalyze poly(ADP-ribosyl)ation reaction to modulate other proteins to maintain chromatin structure, transcription, and genomic integrity (Krishnakumar and Kraus 2010). It is widely known that PARP-1 activation mediates AIF translocation from the mitochondria to nucleus, resulting in caspase-independent cell death (Andrabi et al. 2008; Hong et al. 2004; van Wijk and Hageman 2005). Since the concept of programmed necrosis is accepted, PARP-1-/AIF-mediated caspase-independent cell death has begun to be regarded as necroptosis (Delavallee et al. 2011). However, how the RIP1–RIP3 complex in cytosol signals to PARP-1 in the nucleus is still unclear. In our previous studies, the relationship between PARP-1/AIF and necroptosis by using a hippocampal cell line HT-22 was explored. Glutamate-induced oxidative stress causes a caspase-3-independent necrosis-like cell death in HT-22 cells; the necroptosis-specific inhibitor necrostatin-1 almost completely blocks this kind of cell death (Xu et al. 2007). Necrostatin-1 not only prevents glutamate-induced glutathione depletion in HT-22 cells but also blocks the translocation of AIF from the mitochondria to nucleus (Xu et al. 2007). The effect of necrostain-1 on the AIF upstream signal PARP-1 activity is further examined, and it is found that necrostain-1 inhibits glutamate-induced PARP-1 activation but not total expression. However, necrostain-1 cannot directly inhibit PARP-1 activation induced by the DNA-alkylating agent MNNG (Xu et al. 2010a), indicating that PARP-1-/AIF-mediated signal pathway in HT-22 cell is downstream of necroptosis and that necrostain-1 targets upstream of PARP-1/ AIF, which is consistent with the fact that necrostain-1 inhibits RIP1 activity and blocks the interaction of RIP1 and RIP3 (He et al. 2009). Increasing evidences demonstrate that PARP-1/AIF signal pathway is involved in the downstream signals of necroptosis and may be one of the executioner proteins for necroptosis (Cabon et al. 2012; Deeraksa et al. 2013; Dunai et al. 2012; Jouan-Lanhouet et al. 2012). The mechanisms attending PARP-1/AIF regulation of necroptosis need further investigation.

#### 2.3.4 Mitochondrial Targets

The mitochondrial protein families are involved in the regulation and execution of apoptosis; however, many mitochondrial proteins have been recently found to function in necroptosis. Adenine nucleotide translocase (ANT) and cyclophilin D (CypD), two important components of the mitochondrial permeability transition, regulate the exchange of ADP and ATP between the cytosol and the mitochondria (Halestrap and Brenner 2003). TNF- $\alpha$  and z-VAD-fmk treatment results in a mitochondrial defect in ADP/ATP exchange through the inhibition of ANT activity, which is dependent on RIP1 activation (Temkin et al. 2006). Mice lacking CypD are more resistant to cardiac ischemia and reperfusion injury than control animals and have reduced incidence of necrotic cell death (Nakagawa et al. 2005), suggesting that mitochondrial proteins such as ANT and CypD are important players in necroptosis and that mitochondria functions are also influenced in necroptosis.

Because mitochondria are the major producers of reactive oxygen species (ROS), which are important in executing necroptotic cell death (Festjens et al. 2006b; Moquin and Chan 2010), it is conceivable that RIP1 may directly or indirectly target the mitochondria. Treatment with butylated hydroxyanisole (BHA), a ROS scavenger, inhibits necroptosis in many cell types (Festjens et al. 2006a; Lin et al. 2004). RIP1 and RIP3 act on upstream signals to regulate ROS production during necroptosis. A recent study shows that RIP3 interacts with several mitochondrial enzymes including glycogen phosphorylase (PYGL), glutamate–ammonia ligase (GLUL), and glutamate dehydrogenase 1 (GLUD1) and leads to TNF-/z-VAD-induced ROS production. RIP3 overexpression activates these enzymes and enhances mitochondrial energy metabolism, a process that correlates with enhanced ROS production and necroptosis (Zhang et al. 2009). In contrast, RIP3 deficiency results in reduced ROS production and necroptotic cell death (Cho et al. 2009). Thus, the mitochondria and mitochondrial proteins are important targets for RIP3 or RIP1–RIP3 complex in necroptotic cell death.

#### 2.3.5 JNK/ERK

In a siRNA screen for searching the regulators of necroptosis, transcription factors c-Jun and Sp1 are identified. However, both transcription factors can be activated by JNK signaling (Chuang et al. 2008; Coffey et al. 2002; Hitomi et al. 2008), suggesting that JNK may be downstream of necroptosis (Vanlangenakker et al. 2012). Biton and Ashkenazi recently reported that RIP1 is activated and further

activates NF-κB in the early-phase response upon extensive DNA damage; however, in the absence of caspase activation, RIP1 activates JNK3 pathway in the latephase response, resulting in the activation of the transcription factor AP-1 and FADD-mediated caspase-8 activation (Biton and Ashkenazi 2011). As discussed above, MLKL contains a kinase-like domain which is similar to the mixed lineage kinase that functions as MAPKKs in the JNK pathway (Gallo and Johnson 2002). Therefore, after activation of MLKL by RIP3, MLKL may activate the downstream JNK pathway to execute necroptotic cell death. This is confirmed by Zhao et al.'s findings that MLKL is required for the late-phase activation of JNK in TNF-induced necroptosis in human colon adenocarcinoma HT-29 cells (Zhao et al. 2012). Although current evidences demonstrate the involvement of JNK in the downstream of necroptosis, more studies on its role in necroptosis are needed.

#### 3 Cellular Markers of Necroptosis

For apoptosis and autophagy, many markers were used for the in situ detection in vitro and in vivo, including active caspase-3, cleaved PARP, DNA ladders, Beclin-1 and, microtubule associated protein 1 light chain 3 (Klionsky et al. 2007; Sarkar and Li 2006). Thus far, no specific morphological or biochemical markers are available for the detection of necroptosis. Regardless, many necrotic biomarker candidates and methods for the determination of necroptosis have been proposed recently.

Because the cellular nuclei, mitochondria, lysosomes, and plasma membrane are ruptured upon lethal stimuli on cells, intracellular cytoplasmic contents are released to extracellular spaces. Released proteins, such as high-mobility group box 1 protein (HMGB1), cyclophilin A (CypA), and cytokeratin-18, are regarded as candidate biomarkers of necroptosis (Christofferson and Yuan 2010; Kramer et al. 2004; Scaffidi et al. 2002). HMGB1 protein is a nuclear factor that acts as an architectural chromatin-binding factor, promoting protein assembly on specific DNA targets. When released from the cell, HMGB1 interacts with the receptors for advanced glycation end products and becomes a potent mediator of inflammation, signaling the demise of individual cells to its neighbors (Scaffidi et al. 2002). Since apoptotic cells do not release HMGB1 and fail to promote inflammation (Scaffidi et al. 2002), therefore, released HMGB1 is used to distinguish necrotic cell death from apoptotic cell death. However, its specificity is compromised because HMGB1 can also be actively secreted from immune cells or passively released from cells dying by secondary necrosis following apoptosis (Kazama et al. 2008; Scaffidi et al. 2002). Similar to HMGB1, CypA, a cytosolic peptidyl-prolyl *cis-trans* isomerase, is released in the early stage of necroptosis due to early permeabilization of the plasma membrane (Christofferson and Yuan 2010). Like HMGB1, CypA is also actively secreted; therefore, CypA release may combine with other methods together to determine the existence of necroptosis (Christofferson and Yuan 2010). Intact soluble cytokeratin-18 is released from necrotic cells; but cleaved cytokeratin-18 by caspases is released from apoptotic cells. The ratio of cleaved cytokeratin-18 to

intact cytokeratin-18 in the blood has been proposed as a marker to determine the extent of both types of cell death in patients with endometrial tumors (Kramer et al. 2004). Because not only necrotic cells contribute to intact cytokeratin-18 in the blood, this ratio may provide quantitative and qualitative information in those patients who have a significantly high level of cytokeratin-18 (Linder et al. 2010).

The combination of Tdt-mediated dUTP nick end labeling (TUNEL) staining and cleaved caspase-3 staining has been used recently to determine cell death types in dying cells (Gunther et al. 2011; Welz et al. 2011). Those cells with positive TUNEL staining and positive cleaved caspase-3 staining are considered as apoptotic cells; on the contrary, those cells with positive TUNEL staining and negative cleaved caspase-3 staining are considered as necroptotic cells. Since necroptotic cell death cannot be easily determined using a single method, it should be identified by a combination of methods.

#### 4 Inhibitors and Interventions for Necroptotic Cell Death

Necroptosis is an important cellular death mechanism likely to be involved in human pathology, from viral infections to neurodegenerative disorders. Necroptotic features are commonly observed in a wide range of human pathologies, including stroke, myocardial infarction, brain trauma, and possibly some forms of neurodegeneration. Therefore, inhibition of necroptotic cell death by specific small-molecule inhibitors, necrostatins, may represent an exciting new direction for therapy.

# 4.1 Nec-1 and Its Analogs

Nec-1, a specific inhibitor of necroptosis, acts by inhibiting RIP1 kinase activity (Degterev et al. 2005). Nec-1 has become an important tool for evaluating the contribution of necrotic cell death in vitro and in many experimental animal models in vivo. By using Nec-1, necroptosis has been found in a wide range of cell death paradigms, including ischemic brain injury (Degterev et al. 2005; Xu et al. 2007). Nec-1 shows promising neuroprotection on cerebral ischemia injury for the reduction of infarct volume and the improvement of neurological deficits in middle cerebral artery occlusion (MCAO) mouse models (Degterev et al. 2005; Gu et al. 2010), suggesting that necroptosis is an alternative mechanism to activate delayed ischemic injury, due to its slow-onset kinetics. Nec-1 reduced acute cellular injury, brain tissue damage, motor dysfunction, and spatial learning deficits after controlled cortical impact (CCI) in mice; therefore, Nec-1 and other necrostatins may be promising therapeutic agents to reduce the pathological and functional impairment of traumatic brain injury (You et al. 2008). Moreover, Nec-1 rescues retinal neurons in the inner retinal layers, attenuates retinal degeneration, and reduces functional impairments after retinal ischemic injury (Rosenbaum et al. 2010).

Since the discovery of Nec-1, several structurally distinct Nec-1 analogs for necroptosis blockage have been developed: Nec-3 (Teng et al. 2008), Nec-4

(Teng et al. 2007), Nec-5 (Wang et al. 2007), and Nec-7 (Zheng et al. 2008). Nec-1, Nec-3, Nec-4, and Nec-5 all inhibit RIP1 kinase; however, Nec-7 does not, suggesting that Nec-7 may target a different regulatory pathway from Nec-1.

#### 4.2 Necrosulfonamide

Like Nec-1, necrosulfonamide (molecular weight, 461) – another newly identified specific inhibitor of necroptosis – was selected by screening a chemical library of 200,000 compounds (Sun et al. 2012). Unlike Nec-1 which blocks the interaction of RIP1 and RIP3, necrosulfonamide specifically inhibits necroptosis by targeting MLKL, a key signaling molecule from the mitochondria in the necroptotic cell death pathway, downstream of the active RIP3 (Sun et al. 2012). Necrosulfonamide can block necroptosis at a specific step in different cells such as Jurkat cells, HeLa cells, and mouse 3T3 cells in response to TNF- $\alpha$  stimuli, by binding to the N-terminal of MLKL and covalently modifying MLKL at the reactive amino acid residue cysteine 86 (Sun et al. 2012). Moreover, necrosulfonamide also blocks staurosporine-induced primary necrosis in the presence of a caspase inhibitor in U937 cells (Dunai et al. 2012). Whether necrosulfonamide has inhibitory effects on necroptosis in different disease models in vivo or in vitro is unknown and needs further studies.

# 4.3 Geldanamycin

Because RIP1 has a critical role in RIP1–RIP3 complex formation and RIP1–RIP3dependent necroptosis, RIP1 or RIP3 may be promising new targets for neuroprotection in neurological or other pathological disorders. Our recent study demonstrates that degradation of RIP1 offers an intriguing possibility to rescue necroptotic cells from death induced by oxygen–glucose deprivation (OGD) in the presence of the caspase inhibitor z-VAD (Chen et al. 2012b). RIP1 has low expression in normal cortical neurons, but OGD and z-VAD treatment significantly increases RIP1 expression and induces neuronal necroptosis. However, geldanamycin decreases RIP1 level and rescues OGD- and z-VAD-induced necroptosis (Chen et al. 2012b). Further study indicates that geldanamycin promotes the reduction of heat shock protein (HSP90), which is known to decrease the stability of RIP1 and facilitate the degradation of RIPK1 (Lewis et al. 2000). Therefore, geldanamycin and its analogs may be used for the treatment of stroke or other neurological disorders, by targeting the degradation of RIP1.

# 5 Necroptotic Cell Death in Neurological Diseases

## 5.1 Stroke

Stroke is the third leading cause of death and a leading cause of serious longterm disability. Due to the heavy economic and social burden caused by stroke patients, an understanding of the pathophysiology of stroke is regarded as an urgent and critical research topic in many countries. Recent in vitro and in vivo studies demonstrated that necroptosis is an important process that contributes to neuronal cell death and functional impairment after cerebral ischemic injury (Chen et al. 2012b; Degterev et al. 2005; Meloni et al. 2011; Xu et al. 2010b).

Degterev et al. showed that the specific and potent inhibitor of necroptosis, Nec-1, when administered by intracerebroventricular injection along with the caspase inhibitor z-VAD, significantly reduces infarct volume and improves neurological functions in an MCAO animal model (Degterev et al. 2005). These findings imply that Nec-1 may protect against brain injury by blocking necroptosis when neuronal apoptosis is inhibited by caspase inhibitor z-VAD. Consistent with this, our findings indicate that the combination of apoptosis inhibitor humanin and necroptosis inhibitor Nec-1 provides synergistic protective effects on cerebral infarction and neurological function in mice (Xu et al. 2006; Xu et al. 2010b). By using an in vitro OGD ischemia model, a recent study demonstrated RIP1-dependent necroptosis induced by z-VAD treatment; geldanamycin, which promotes the degradation of RIP1, rescues OGD- and z-VAD-induced necroptotic cell death in culture cortical neurons (Chen et al. 2012b). A recent study with cell death biomarkers also identified that different forms of cell death including necroptosis exist in both acute and delayed OGD models (Meloni et al. 2011). These studies imply that necroptosis is a potential therapeutic target for the treatment of stroke.

#### 5.2 Neonatal Hypoxic–Ischemic Encephalopathy (HIE)

Neonatal HIE is the most common cause of perinatal brain injury, resulting from pulmonary and/or cardiac dysfunction. Because of the similarity, rodent neonatal hypoxia-ischemia (HI) injury models are used to study the mechanisms of HIE (Ashwal and Pearce 2001). West et al. (West et al. 2006) demonstrates that caspase-3-deficient mice have greater tissue loss than controls after cerebral HI injury; caspase-3 knockout results in increased PARP-1-mediated caspase-independent cell death following HI. By using the same animal model, Northington et al. (Northington et al. 2011) found that Nec-1 significantly decreases brain injury in the forebrain and thalamus and decreases necrotic cell death, suggesting the existence of necroptosis in HI brain injury. They also showed that HI brain injury promotes necroptosis by increasing RIP1-RIP3 complex formation and inhibiting RIP3-FADD interaction and that mitochondrial dysfunction in neurons and astrocytes following neonatal hypoxia-ischemia is prevented by RIP1 activity inhibition (Chavez-Valdez et al. 2012; Northington et al. 2011). Therefore, the protection of necrostatin-1 on HI brain injury may represent a therapeutic strategy and new direction for therapy of neonatal HI.

#### 5.3 Trauma

A recent study also demonstrates that RIP1-/RIP3-dependent necrosis appears to be involved in the mechanism of cell death and neurological dysfunction after traumatic brain injury (You et al. 2008). Necrostatin-1 administration significantly reduces brain lesion volume and motor deficits after controlled cortical impact in a trauma animal model but also improves spatial memory acquisition by Morris water maze performance. However, Nec-1 did not reduce caspase-3-positive cells in dentate gyrus or cortex, indicating that Nec-1 is not an apoptosis inhibitor, but a necroptosis inhibitor (You et al. 2008).

#### 5.4 Huntington's Disease

Huntington's disease is a neurodegenerative genetic disorder characterized by movement abnormalities, psychiatric symptoms, and cognitive deficits. Zhu et al. (Zhu et al. 2011) recently examined the role of necroptosis in Huntington's disease by using Nec-1 as tool in immortalized striatal ST14A cells and R6/2 transgenic mice. In ST14A cells stably expressing mutant Htt fragment (N548-128Q), a cell model of Huntington's disease, the pancaspase inhibitor z-VAD induces caspase-3-independent cell death. Caspase-8-specific inhibitor (IETD-fmk), but not caspase-3- and caspase-9-specific inhibitor, induces a similar cell death as z-VAD in ST14A cells. Nec-1 almost completely rescues this kind of cell death and inhibits RIP1 cleavage induced by z-VAD or caspase-8 inhibitor IETD-fmk, supporting the notion that Nec-1 blocks RIP1-dependent necroptosis in ST14A cells (Zhu et al. 2011). In the R6/2 transgenic mouse model of Huntington's disease, Nec-1 treatment significantly delays the onset of Huntington's disease as well as behavioral deterioration and extends the life span, suggesting the involvement of necroptosis in the pathogenesis of Huntington's disease (Zhu et al. 2011). This study implies that necroptosis may be involved in other neurodegenerative disorders such as Parkinson's disease and Alzheimer's disease. Moreover, Nec-1 may be a promising protector in these diseases.

### 5.5 Glutamate Toxicity

Glutamate excitotoxicity is linked to neurological disorders such as stroke, Parkinson's disease, Huntington's disease, and Alzheimer's dementia (Beal 1992; Lau and Tymianski 2010). Glutamate excitotoxicity also causes neuronal death in cultured primary neurons in vitro (Chen et al. 2012a; Zhang et al. 2011b). Nec-1, as a necroptosis inhibitor and in the absence of caspase inhibitors, protects against glutamate analog *N*-methyl-D-aspartate (NMDA)-induced cell death in primary cortical neurons (Li et al. 2008). In this study, Nec-1 partially blocked NMDA effects, suggesting necroptotic involvement in NMDA-induced cell death. However, necroptosis only occupies a small portion of cell death without the induction of caspase inhibitors. Non-receptor-mediated glutamate toxicity causes necroptosis in neuronal cells, as well (Xu et al. 2007; Xu et al. 2010a). In a hippocampal HT-22 cell line that has no ionotropic glutamate receptor, again, glutamate produces caspase-3-independent cell death. Nec-1 almost completely blocks glutamate-induced cell death, suggesting the involvement of necroptosis in glutamate-induced cell death in HT-22 cells (Xu et al. 2007). Further study indicates that the PARP-1/AIF pathway mediates necroptosis induced by glutamate in HT-22 cells (Xu et al. 2007; Xu et al. 2010a). These in vitro studies imply that glutamate toxicity may cause necroptosis in many neurological disorders, thus providing a new direction for treatments.

#### 6 Conclusion

Necroptosis is a recently identified regulated cell death that has the morphological features of necrosis. Investigation into the regulation and signaling pathways of necroptosis represents one of the hot topics in cell death research. Recent findings throw light on the importance of necroptotic cell death in normal development, immunity, inflammation, cancer, and neurological disorders (Han et al. 2011; Vanlangenakker et al. 2012; Yuan and Kroemer 2010). An explosion of recent articles begins to unravel the mysteries centering on the necroptotic process. In response to external or internal stimuli, RIP1 and/or RIP3 are the determining factors at the central of the regulation platform of necroptosis to function as the switches between apoptosis to necroptosis. Pro-apoptotic proteins such as caspase-8, FLIPL, and FADD negatively regulate RIP1-RIP3 complex formation and inhibit the process of necroptosis. However, some factors such as CYLD, which deubiquitinates or modifies RIP1 and induces RIP1-RIP3 complex formation, positively promote necroptotic cell death. Although the complete signaling pathway remains unclear, recent advances have shown that mitochondrial enzymes MLKL and PGAM5 are directly downstream from RIP3 activation and that DRP1 may be one of the executioners of necroptosis. Due to complicated signaling pathways in the cells, more investigations are needed to remove the veil over this cell death process. The unraveling of the mechanisms governing necropototic cell death will lead to a better understanding of the interplay of necroptosis and other forms of cell death and the identification of new clinical targets.

Because of the lack of clear and distinctive biomarkers for necroptosis, it is difficult to determine the existence of necroptosis in animal disease models. In addition, it is hard to tell whether necroptosis is the only form of necrotic cell death. More specifically biomarkers need to be developed to effectively discern necroptosis in vivo. Thus far, only Nec-1 and NSA are considered as specific inhibitors for necroptosis – used as tools to examine necroptosis. Because these inhibitors may also block other cell death processes, it is possible that there is an overestimate of the portion of cell death deemed to be necroptosis.

Recent studies have examined the existence of necroptotic cell death in many neurological disorders such as stroke, trauma, and Huntington's disease. Because no specific biomarker is available, most studies used the neuroprotection of Nec-1

on cell death as the evidence for necroptosis. In addition, it remains difficult to understand the contribution to neurological pathogenesis; how great a percentage of necroptosis occurring in the pathology of neurological disorders is still difficult to estimate. The extensive studies on necroptosis hopefully will lead to the identification of specific necrotic biomarkers to resolve the many questions. Because caspase inhibitors promote necroptosis in vitro, does necroptosis play a critical role in the pathology of neurological disorders under the circumstances wherein there is no caspase inhibition? This remains an unanswered question. Recent studies demonstrate that mitochondrial dysfunction is involved in the mechanisms of necroptosis (Sun et al. 2012; Wang et al. 2012; Zhang et al. 2009; Zhao et al. 2012) and that mitochondrial dysfunction contributes to the pathophysiology of acute neurological disorders such as stroke and neurodegenerative disorders such as Parkinson's disease (Perez-Pinzon et al. 2012; Schon and Przedborski 2011; Vosler et al. 2009). Therefore, the connection between necroptosis and neuronal cell death in these disorders is in need of further investigation. Because of its potential involvement in neurological disorders, necroptosis has become a potential thera-

peutic target for these disorders. A clearer understanding of necroptotic mechanisms will provide new targets and exciting opportunities to develop more potent neuroprotective agents such as Nec-1 or necrosulfonamide for clinical treatments of neurological disorders.

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# **RCSN Cell System for Identifying Dopaminergic Neurotoxicity**

# Pablo Caviedes, Raúl Caviedes, and Juan Segura-Aguilar

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#### Abstract

Dopaminergic toxicity represents a potential mechanism underlying Parkinson's disease (PD) neuropathology. Nevertheless, the study of such a mechanism is hampered by the lack of permanent and stable in vitro models that bear relevant cellular traits, namely, neuronal dopaminergic function. Although various permanent cell lines exhibiting variable dopaminergic properties do exist, such properties are not necessarily stable and may require the application of complex and costly differentiation protocols for induction. The latter is particularly true when inducing in vitro differentiation from more undifferentiated tissue, such as stem cells. Also, cell lines may lose viability or eventually undergo permanent differentiation. This chapter discusses

P. Caviedes (🖂) • R. Caviedes • J. Segura-Aguilar

Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

e-mail: pcaviede@med.uchile.cl

a permanently growing cell line, named RCSN-3, which was established from the substantia nigra of an adult Fisher 344 rat. The cell line retains dopaminergic traits, including dopamine production and secretion, and the presence of catecholamine reuptake transporters. Notably, these properties have remained expressed in RCSN-3 cells for decades. This chapter also addresses the contribution of RCSN-3 to dopaminergic-mediated toxic phenomena, in particular, and to the identification of potential therapeutical targets in dopaminergic neurons. Finally, RCSN-3 cells are also presented as a model for cell transplant therapy, in animal models of PD, and their contribution in this field is discussed in relation to more recently available cell sources, such as stem cells and induced pluripotent stem cells (iPSCs).

# 1 Neurons in Culture: What Can They Do for Us?

The culture of neuronal cells has been widely used, as it presents an unparalleled opportunity to provide model systems for ready and unrestricted access to living tissue. Neurons, however, exhibit quite distinct features when taken from an in vivo to in vitro environments, which must be addressed to provide adequate answers for the questions to be asked. Indeed, neuronal tissue inherently has greater demands (i.e., oxygen, nutrients, trophic factors), which must be provided to insure adequate survival and maintenance of differentiated function in culture conditions (Freshney 2000). Among them, one can count the use of special matrixes for attachment and differentiation (collagen, laminin, fibronectin, extracellular matrix preparations, etc.), along with the supplementation of soluble factors in the media, which have a very wide range of complexity, ranging from monovalent ions to molecules of high molecular weight (Levi-Montalcini 1979; Marchionni 1993). These demands, associated with limitations in the quantities of procurable tissue, particularly from human origin, have prompted the interest in developing immortalized cell lines from neurons, which would potentially provide a limitless amount of cells and, hopefully, stable models for years to come. However, many such cell lines tend to exhibit problems in their stability and/or viability, with end results ranging from death to irreversible loss of differentiated functions (Cárdenas et al. 1999). Yet, if such drawbacks can be overcome, they appear as a feasible alternative, as they can generate genetically homogeneous cell populations that can be suitable models to explore physiological and pathophysiological phenomena at the cell level and also identify pharmacological targets which can be used for drug screening.

Following this line of thought, the use of tissue culture in dopaminergic systems appears as an adequate tool to study problems of clinical relevance, especially those related to relevant degenerative diseases such as Parkinson's disease (PD). Indeed, the discrete localization of the neuropathological hallmark

in this condition – namely, the degeneration of the *substantia nigra pars compacta* (SN) – presents dopaminergic culture systems as ideal models for the study of PD cell pathophysiology. This chapter will refer to such models, and it will particularly address a specific neuronal cell line, named RCSN-3, which was established from the substantia nigra of an adult Fisher 344 rat. Evidence of its use in the study of potential toxic mechanisms relevant to PD will be discussed, along with future directions.

# 2 Parkinson's Disease: Where We Stand...

Parkinson's disease is a neurodegenerative disorder affecting an estimated 0.3 % of the population in the industrialized world (de Lau and Breteler 2006), the elderly being the highest population at risk. The disease is progressive in nature, and there are no effective treatments to arrest or revert the evolution of the disease. The clinical manifestations of PD are several, and presentations vary greatly among patients. The most evident include signs and symptoms that affect voluntary movement, which include resting tremor in the extremities and muscular rigidity, which leads to difficulties in walking with a characteristic shuffling gait. Also, impaired writing, speaking, and masking of facial expression are present. Bradykinesia, or slowness in initiating and executing movements, and stooped posture and instability are also commonly observed signs (Sian et al. 1999). Neuropsychiatric manifestations (mood disorders, altered thought process, cognitive decline) may also be present (Jankovic 2008), particularly in later stages of the disease, thus presenting PD as a far more complex condition than one that merely compromises motor function. At the neuropathological level, the most remarkable finding is the degeneration of the dopaminergic neurons present in the SN. These neurons project to the striatum, providing the latter with a basal dopaminergic input necessary for the adequate function of the regulatory circuit of the basal nuclei. The degeneration of the SN determines an imbalance in the regulation of the circuitry of the basal ganglia, whose GABAergic output to the thalamus is increased, resulting in an overall inhibition of this projection nucleus (Lozano et al. 1998). Hence, messages processed in the basal ganglia which project to the cerebral cortex via the thalamus have an increased inhibitory component in PD, especially those coding for voluntary movement, resulting in reduced motility.

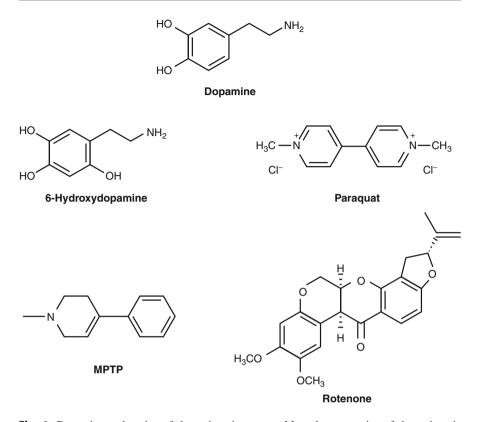
The mechanisms for the progressive degeneration of the SN in PD remain largely unknown, in particular for sporadic forms of PD. A small minority of patients with mutations in certain genes such as alpha-synuclein, parkin, leucine-rich repeat kinase 2 (LRRK2), or PTEN-induced putative kinase 1 (PINK1) will develop the disease with a high degree of certainty (Davie 2008; Lesage and Brice 2009). Yet, the mechanisms of toxicity remain obscure, in particular at the cellular and molecular level. Therefore, the development of neuronal in vitro cell models

that resemble SN function is of the essence in order to address such questions and in turn identify relevant intracellular therapeutical targets.

#### 3 Ex Vivo Dopaminergic Cells: What can They do to Help?

Many strategies are being pursued to develop new therapies for PD patients, mainly targeting to the prevention of the damage of the nigrostriatal system (neuroprotection or rescue) and/or to replacing neurons that were lost due to the underlying condition. Efforts in this regard have included the use of trophic factors (Takayama et al. 1995) and viral vectors (Choi-Lundberg et al. 1997) for protection and rescue, to the transplantation of primary xenogeneic or allogeneic tissue for repair (Deacon et al. 1997). Nevertheless, progress in the knowledge of mechanisms is hampered, firstly, by the lack of animal models that adequately reflect the human condition. Indeed, PD does not occur naturally in other mammals, and most animal models involve acute destruction of the SN by use of toxins, most of which share structural resemblance to dopamine (DA) (i.e., 6-OH-dopamine intoxicated rats; 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP)-intoxicated monkeys) (see Fig. 1). Hence, such models do not reproduce the inherent chronic damage of the SN in PD and tend to target a reduced number of many potential mechanisms involved in dopaminergic degeneration. Indeed, such toxins result in rapid and sizable oxidative stress and/or energy depletion, which are most probably not solely responsible for cell damage. Other models have incorporated transgenics for known PD-related genes, including knockouts of alpha-synuclein (SNCA) or overexpression of mutated forms of this protein (Blandini and Armentero 2012). In spite of such progress, no such models reproduce all the characteristics of PD, and the aforementioned models tend to address specific questions of a given mechanism.

The first approach to a given condition must address alterations at the cellular level, a goal that is difficult to attain in a full animal, where many variables escape from the control of the researcher. Initially, in vitro models comprised primary cultured cells, which tend to reflect more closely the traits of the tissue of origin. Yet, the inherent variability of such preparations and limitations in the amounts of tissue procurable conspire against presenting these models as a universal preparation. The use of neuronal cell lines overcame the noted limitations in quantities of tissue, yet longterm instability and eventual permanent dedifferentiation are but two factors that can hinder their usefulness. Nevertheless, dopaminergic cell lines have been used for quite some time. The PC12 cell line, which is derived from the pheochromocytoma of a rat (Greene and Tischler 1976), has been and is still largely used to study dopaminergic and neuronal function, yet it is tumorigenic, it is difficult to grow, and it requires nerve growth factor (NGF) for neuronal differentiation in vitro. Another interesting cell line, SH-SY5Y, derived from the neuroblastoma of a child, has also been used to study dopaminergic function, especially since it has dopamine- $\beta$ -hydroxylase activity (Oyarce and Fleming 1991), thus suggesting the presence of catecholaminergic function, yet it does not derive from a dopaminergic precursor.



**Fig. 1** Dopamine and toxins of dopaminergic system. Many known toxins of dopaminergic neurons share structural similarities with dopamine, thus entering dopaminergic neurons via dopaminergic reuptake transporters. 6-Hydroxydopamine generates reactive oxygen species such as superoxide radical inside the cells. MPTP (1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine) is the precursor of the neurotoxin MPP<sup>+</sup> (1-methyl-4-phenylpyridinium), which, once inside dopaminergic neurons, interferes with complex I of the electron transport chain. Rotenone interferes with the electron transport chain in mitochondria, particularly by inhibiting the transfer of electrons from iron-sulfur centers in complex I to ubiquinone, in turn interfering with NADH in the creation of ATP. Paraquat, a known herbicide, can catalyze the formation of reactive oxygen species (ROS), particularly superoxide

Lately, stem cells, derived from the nervous system (neural stem cells), have been used to generate sufficient cell mass to later differentiate to dopaminergic cells. Yet, the reduced amount of cells, slow in vitro proliferation, and the ethical and practical difficulties involved in procuring viable human nervous tissue do introduce difficulties in this line of work. At present, one of the most interesting lines of research involves reprogramming human fibroblasts from patients to generate induced pluripotent stem cells or iPSCs. Such cells can generate patient-specific cells that can later be differentiated to dopaminergic neurons (Swistowski et al. 2010).

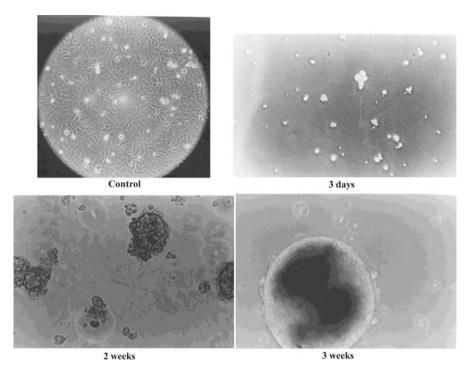
# 4 RCSN-3: A Permanent In Vitro Neuronal Dopaminergic Model

All of the aforementioned cell models are and have been useful in the study of the neuronal physiology and pathophysiology of PD. Yet, limitations in growth rate, in differentiation, and in the source of tissue define most of their limitations, whereas lack of stable, relevant dopaminergic functional traits is another drawback. Hence, a stable dopaminergic neuronal cell line, derived from SN tissue that retains differentiated traits, would provide a valuable model to test PD-related phenomena at the cell level. In this regard, RCSN-3 appears as an interesting candidate for such tasks.

RCSN-3 was established in the early 1980s, from primary cultures of the SN dissected from a 4-month-old Fisher 344 rat (Arriagada et al. 2002). The cell line was established from the said primary cultures by means of the UCHT1 immortalization protocol (Caviedes et al. 1993, 1994; Cárdenas et al. 1999, 2002). This original procedure has managed to successfully generate neuronal cell lines in permanent culture from diverse origins, obtained from both normal and pathological subjects (Caviedes et al. 1993; Liberona et al. 1997; Allen et al. 2000). Briefly, the immortalization procedure entails maintaining neuronal primary cultures in the presence of media conditioned by the rat thyroid cell line UCHT1. This treatment induces tissue immortalization in variable periods of time (Caviedes et al. 1993). This procedure has allowed the establishment of continuously growing cell lines that express stable neuronal phenotypes and lack glial characteristics. These and other lines so established retain differentiated traits, some for decades (Caviedes et al. 1993; Liberona et al. 1997; Allen et al. 2000) and 2005).

After a period of 7–8 months of UCHT1 treatment, transformation foci of rapid proliferation became evident, after which the cultures were weaned from UCHT1 media. The resulting motherline, named RCSN, was subsequently cloned. Of these, RCSN clone 3, or RCSN-3, has been the line that stably retains dopaminergic traits. The RCSN-3 cell line grows in monolayers and can also be induced to grow in suspended, attachment-free clusters (Fig. 2) (Caviedes et al. 2012). RCSN-3 has a doubling time of approximately 50 h, a plating efficiency of 21 %, and a saturation density of 50,000 cells/cm<sup>2</sup> when grown in basal growth media composed of DMEM/HAM-F12 nutrient mixture (1:1), 10 % adult bovine serum, 2.5–5 % fetal bovine serum, and 40 mg/l gentamicin (Dagnino-Subiabre et al. 2000; Paris et al. 2001; Martinez-Alvarado et al. 2001).

Regarding their neuronal trails, RCSN-3 cells are immunohistochemically positive for neuron-specific enolase (NSE), MAP-2, 200 kD neurofilament, and parvalbumin. Further, RCSN-3 cells exhibit intracellular melanin deposits, as evidenced by Fe<sup>2+</sup> ion capture cytology studies (Arriagada et al. 2002). On the other hand, glial markers such as GFAP and S-100 are negative (Arriagada et al. 2002). Regarding dopaminergic function, the cells express tyrosine hydroxylase (TH) in basal culture conditions, with increased labeling after differentiation with serum deprivation (Arriagada et al. 2002; Paris et al. 2008). They also



**Fig. 2** RCSN-3 cells grow in monolayers when kept in standard growth media. When subjected to culture protocols that disfavor attachment (Caviedes et al. 2012), RCSN-3 cells grow in suspension and cluster progressively

show fluorescent labeling in the cytoplasm after treatment with paraformaldehydeglyoxylic acid, suggesting the presence of catecholamine content. Experimental evidence of active DA release by RCSN-3 cells has been obtained with HPLC studies of culture supernatant and single cell amperometry with carbon electrodes, both of which are highly suggestive that RCSN-3 is capable of secreting this neurotransmitter. Further, RCSN-3 cells express a number of transporters, such as norepinephrine, serotonin, and, interestingly, DA (Paris et al. 2008). Hence, this is the only known dopaminergic cell line that originated from the SN of an adult animal and thus an attractive model to study PD-related mechanisms, in particular those related to dopaminergic toxicity.

#### 5 Can Dopamine be Harmful? How?

Firstly, the causes for the discrete compromise of dopaminergic neurons of the SN in PD patients remain a mystery. Yet, DA itself may have a role. As a catecholamine, DA has a catechol ring with two hydroxyl groups in positions 4 and 5 (Fig. 1). In physiological conditions, such hydroxyl groups may remain

reduced due to several mechanisms, such as inclusion in low-pH secretory vesicles, reduction by several scavenging enzymes, and formation of neuromelanin. An imbalance in such mechanisms could shift the balance of the redox state of such hydroxyl groups to a more oxidative stage – namely, superoxide – this introducing a potential pro-oxidative state. Considering the specificity of the lesions in PD, it is possible then that the high oxidative state associated with DA metabolism may cause deterioration of dopaminergic neurons. The mechanism underlying increased oxidative stress may involve DA itself, because oxidation of cytosolic levodopa/DA may be deleterious to neurons. Indeed, DA causes apoptotic cell death as shown by morphological nuclear changes and DNA fragmentation (Luo et al. 1998). DA and its metabolites seem to exert cytotoxicity mainly by generating highly reactive quinones through auto-oxidation (Martinez-Alvarado et al. 2001). On the other hand, the toxicity of DA may be due to the generation of reactive oxygen species (ROS) that could disrupt cellular integrity, causing cell death. Indeed, aberrant regulation of DA is accompanied by accumulation of DA metabolites that could induce toxicity (Arriagada et al. 2004; Hattori et al. 2009).

The possibility that environmental factors may play a role in sporadic PD was addressed over a decade ago (Tanner et al. 2009). Epidemiological risk factor analyses of typical PD cases have identified several neurotoxins, including the aforementioned active metabolite of MPTP, 1-methyl-4-phenyl-1,2,3,6tetrahydropyridinium ion (MPP<sup>+</sup>), paraquat, dieldrin, manganese, copper, and salsolinol (Chun et al. 2001; Paris et al. 2008). Regarding MPP<sup>+</sup>, Chun et al. (2001), using a nigral dopaminergic cell line, SN4741, demonstrated that such neurotoxins can induce apoptotic cell death through a common molecular mechanism. These authors observed initial increases in H<sub>2</sub>O<sub>2</sub>-related ROS activity, followed by activation of JNK1/2 MAP kinases; activation of the PITSLRE kinase, p110, by both caspase-1- and caspase-3-like activities; and, finally, apoptotic cell death. Pharmacological intervention with combinations of the antioxidants significantly protected the cells against such damage.

The SN4741 model would then appear as an interesting model to study PD-relevant pathophysiology. Yet, this line was established from transgenic mouse embryos containing the targeted expression of the thermolabile SV40Tag in SN neurons. Cell lines generated in this fashion tend to show instability and loss of function (Frederiksen et al. 1996), a problem not reported for UCHT1-established cell lines. Further, SN4741 cells are embryonic in origin and require differentiation in vitro. In this regard, RCSN-3 cells express dopaminergic traits without need of lengthy or costly differentiation protocols, and it has retained such traits for years. Interestingly, they also undergo toxicity and death upon exposure to such the aforementioned neurotoxins (Paris et al. 2008). Also, the RCSN-3 cell line expresses the reuptake transporters for DA (DAT) and norepinephrine (NET) (Paris et al. 2005). These properties of RCSN-3 cells allowed the discovery of a novel mechanism involving heavy metal toxicity (Cu<sup>2+</sup>, Fe<sup>2+</sup>) related to formation of complexes with dopamine (Paris et al. 2001, 2005); and their subsequent entry into the cell specifically by these transporters has provided new

light on how such environmental factors can contribute to PD pathogenicity. Indeed, by using highly efficient transport systems, heavy metals can then enter rapidly, generating oxidative stress in submitochondrial compartments via the Fenton reaction (Thomas et al. 2009).

Inside the cell, DA can induce toxicity, in particular when its redox state is shifted toward more pro-oxidizing conditions. Indeed, DA-oxidized metabolites can inhibit the respiratory chain in mitochondria (Ben-Shachar et al. 2004), thus producing an energy crisis in the cell. In this regard, RCSN-3 cell can undergo toxicity phenomena with treatment of DA oxidation products such as aminochrome treatment, once reductases such as DT-diaphorase are inhibited (Arriagada et al. 2000, 2004). RCSN-3 cells also express the vesicular monoamine transporter-2 – VMAT-2 (Fuentes-Bravo et al. 2007). When this transporter was selectively inhibited with reserpine, exposure of cell to external DA induced clear apoptotic phenomena, such as blebbing and chromatin condensation. Also, mitochondrial damage was evident at the electron microscopy level (Fuentes-Bravo et al. 2007).

#### 6 Cell Transplantation in PD: The Final Frontier?

Cell transplantation therapy was proposed several decades ago as a curative therapy for PD. The rationale for such an approach was based on the discrete anatomical location of the neuronal population compromised and the fact that dopaminergic output from the SN to the striatum has a rather tonic form of discharge. Hence, replacing such tissue with new, DA-secreting cells appeared as clinically promising experimental treatments, in particular in late-stage PD, where tissue replacement is essential. Efforts so far have reached the clinical setting in PD and other neuropathological conditions, using sources such as fetal tissue (Olanow et al. 2006), xenografts from porcine tissue (Borlongan and Sanberg 2002; Poncelet et al. 2009), and immortalized cell lines (Prasad et al. 1998; Paul et al. 2007; Miljan et al. 2009). Indeed, considering that PD patients develop symptoms when approximately 50-70 % of the dopaminergic neurons in the SN are lost (Lozano et al. 1998), the development of cell transplant models appears as a necessary step in the cure for the disease. This is also relevant in light of the lack of predictive markers of the disease; hence, diagnosis is clinical and essentially tardy. Further, the discrete anatomical localization of the degenerating tissue and the large size of the target nucleus (striatum) make it a likely candidate for cell transplant therapy with minimally invasive stereotaxic procedures. In this regard, dopaminergic cell lines can provide a model for testing such therapeutical approaches, considering the existence of relevant animals models such as the 6-OHDA rat and MPTP-intoxicated monkeys. Cell lines are easy to manipulate, and genetic modifications set forth to improve dopaminergic function and/or life expectancy of the graft (i.e., transfection with genes that encode for neurotrophic factors) may provide pertinent, useful information to determine conditions that optimize these functions in transplantable dopaminergic cells.

# 7 Cell Therapy: RCSN-3 Contribution and the Use of Induced Pluripotent Stem Cells (iPSCs)

RCSN-3 cells have also been used as in cell transplant approaches, using induced PD animal models. Indeed, implant of RCSN-3 in the striatum of 6-OHDA-intoxicated rats resulted in a stable reduction of around 50 % in apomorphine-induced rotations. Lately, the development of methodologies to generate pluripotent stem cells from somatic tissues (iPSCs) has provided researchers with an interesting source of stem cells, with the important added value of being patient specific. Hence, the first advantage of iPSC technology is the generation of relevant in vitro models for diseases and drug treatment assays (Soldner et al. 2009). Further, protocols to increase the yield of dopaminergic neuron generation from iPSC from either humans or mice are now available (Park et al. 2008; Soldner et al. 2009; Hargus et al. 2010; Devine et al. 2011). Indeed, neural precursor cells and dopaminergic neurons have been generated from healthy iPSC (Wernig et al. 2008), which when transplanted in the brain can integrate in the parenchyma, migrate, differentiate, and display neuronal function such as action potentials. Most interestingly, grafts derived from iPSC are capable of restoring motor function in animal models for PD (Wernig et al. 2008), suggesting that dopaminergic neurons derived from iPSC can survive implantation and remain functional in vivo.

In humans, iPSC-derived dopaminergic neurons have been obtained from healthy donors (Swistowski et al. 2010), sporadic PD patients (Soldner et al. 2009; Hargus et al. 2010), or PD patients bearing some of the known mutations (Devine et al. 2011; Martínez-Morales and Liste 2012). For instance, DA neurons can be generated from iPSCs that carry a mutation in *LRRK2* gene (p.G2019S), the most common PD-related mutation (Drolet et al. 2011). After reprogramming, and applying a culture differentiation protocol using feeders, iPSC-p.G2019S generated significant amounts of DA. This constitutes the most interesting application for such models, as "custom" therapies can potentially derive from such unique models.

#### 8 Conclusion

Overcoming neurodegenerative diseases, of which PD represents an important fraction, is the most demanding challenge. Relevant therapies will come from the understanding of cellular and molecular mechanisms involved in cell damage, along with the development of predictive markers. The lack of the latter introduces yet another limitation in the quest for new therapies, where neuroprotective strategies are clearly at a loss.

Relevant cell models are a must in the undertaking of both the aforementioned tasks. Novel treatments require models such as RCSN-3 cells for first-level screening and may also play a part in the study of predictive markers. The development of iPSCs is the most exciting chapter in the field, as they can provide disease-specific models from patients. Eventually, they may constitute adequate

sources for cell transplant therapy. Yet, much is to be said about the long-term stability of iPSCs; hence, their routine application to cell transplant therapy may be further away than desired, as well as their application as in vitro models. Until that day, well-established models like RCSN-3 will have a role in addressing the problem at hand.

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# Microglia: Neuroprotective and Neurodestructive Properties

## G. Jean Harry and Christopher A. McPherson

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### Abstract

In the central nervous system, resident macrophages, microglia, and perivascular macrophages perform similar functions to peripheral macrophages yet display highly specialized features. They maintain some phenotypic characteristics and lineage-related properties common to their cells of origin and their rapid response in areas of neuronal death led to the contention that microglia serve as brain macrophages. While microglia are often considered the immune cell of the brain, they also show distinct features that make them unique from other tissue macrophages. They are maintained in a relatively quiescent and monitoring state by regulatory factors released by neurons and astrocytes. Via contact-dependent and receptor-dependent signaling, microglia rapidly respond to various events in the brain in an effort to return the microenvironment to homeostasis, to assist in refining the neural network

G.J. Harry (🖂) • C.A. McPherson

Neurotoxicology Group, National Toxicology Program Laboratory, National Institute of Environmental Health Sciences, Research Triangle Park, NC, USA e-mail: harry@niehs.nih.gov; mcphers1@niehs.nih.gov

during development and with repair, and to remove excess and aberrant proteins. Depending on the nature of the response and the physiological function of the microglia, a release of immune-related signaling factors can accompany a morphological change. It was initially thought that the immune-related response of microglia mirrors that of peripheral macrophages and considered to be adverse to the nervous system. However, further evaluation of these cells suggests that overall their responses are ones critical to the maintenance and function of the nervous system. This chapter will serve to set the framework for evaluating the microglia within the context of data available on their varied functions within the nervous system.

#### **Keywords**

Cytokines • Microglia • Neurodegeneration • Neuroinflammation • Pattern recognition receptors

#### List of Abbreviations

	eviations
AD	Alzheimer's disease
AMPA	Alpha-amino-3-hydroxy-5-methylisoxazole-4-propionic acid
ApoE	Apolipoprotein
ATP	Adenosine triphosphate
$A\beta_{42}$	Amyloid-beta 42
BBB	Blood–brain barrier
Ca <sup>2+</sup>	Calcium
CNS	Central nervous system
COX	Cyclooxygenase
CR	Complement receptor
CSF	Colony-stimulating factor
CX3CL1	Fractalkine
DAMPs	Danger-associated molecular patterns
Fc	Fragment, crystallizable region
HO-1	Heme oxygenase-1
HSP	Heat shock protein
IFN	Interferon
IL	Interleukin
iNOS	Inducible nitric oxide synthase
LPS	Lipopolysaccharide
mGluR	Metabotropic glutamate receptor
MHC	Major histocompatibility complex
MMP	Matrix metalloproteinase
MPP+	1-methyl-4-phenylpyridinium
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine
NF-κB	Nuclear factor kappa B
NLRs	NOD-like receptors
NO	Nitric oxide

NOS NOX2 P2 PAMPs PD PRRs RAGE RNS ROS SN	Nitric oxide synthase NADPH oxidase Purinergic Pathogen-associated molecular patterns Parkinson's disease Pattern recognition receptors Receptor for advanced glycation end products Nitric oxide-dependent reactive nitrogen species Reactive oxygen species Substantia nigra
PD	Parkinson's disease
PRRs	Pattern recognition receptors
RAGE	Receptor for advanced glycation end products
RNS	Nitric oxide-dependent reactive nitrogen species
ROS	Reactive oxygen species
SN	Substantia nigra
TGF	Transforming growth factor
TLR	Toll-like receptor
TNF	Tumor necrosis factor
TREM-2	Triggering receptor expressed on myeloid cells-2

### 1 Introduction

Macrophages are phagocytic cells originating from precursor cells in the bone marrow that are transported via the peripheral blood as monocytes to become tissue-specific macrophages (Davoust et al. 2008; Ginhoux et al. 2010; Morris et al. 1991). These cells maintain a first line of defense against injury and infection to return the system back to homeostasis by recruiting various effector molecules and other immune cells to assist in the task of killing pathogens and restoring tissue integrity (Gordon and Martinez 2010; Mosser and Edwards 2008). Two arms of the immune system, a rapid innate immune response and a slow adaptive immune response, are responsible for these functions (Lo et al. 1999). Innate immune receptors are preformed and stably encoded in the genome, while those of the adaptive immune system are both stochastically generated and shaped by the types and frequency of pathogens the cell encounters. First-responder cells rapidly recognize and respond to cell damage or infection-related dangerassociated molecular patterns (DAMPs) increasing phagocytic activity. They also respond to pathogen-associated molecular patterns (PAMPs) expressed by pathogens (e.g., lipopolysaccharide (LPS)) but not by host cells. In responding, there is a production of cytotoxic reactive oxygen species (ROS), nitric oxide (NO), and proinflammatory cytokines such as tumor necrosis factor (TNF), interleukin (IL)-1, and IL-12. The second arm provides adaptive immunity to detect and respond to novel as well as evolutionarily conserved pathogenic molecules, i.e., antigens (Lo et al. 1999; Schwartz and Kipnis 2011). The adaptive immune system is comprised of long-lived lymphocytes (T cells and B cells) and antigen-presenting cells. The antigen-driven expansion of clonal populations of lymphocytes allows for experience to shape the response and for establishment of memory lymphocytes that are able to more rapidly attack a reencountered antigen.

Under such conditions, antigen-presenting cells play a critical role by phagocytizing material, loading the captured antigen into the pocket of the major histocompatibility complex (MHC) molecule for cell-surface antigen presentation on B cells, macrophages, and dendritic cells to facilitate recognition and clearance (Lo et al. 1999).

In the central nervous system (CNS), resident macrophages, microglia, and perivascular macrophages perform similar functions to peripheral macrophages yet display highly specialized features. In a temporal and spatial manner, monocyte colonization of the brain occurs with tissue vascularization and contact with vascular sprouts prior to birth (Cuadros and Navascues 1998; Marin-Padilla 1985; Rymo et al. 2011). The resulting microglia are yolk-sac derived cells maintained independently of blood-borne hematopoietic progenitors (Ajami et al. 2007; Ginhoux et al. 2010). They maintain some phenotypic characteristics and lineagerelated properties common to macrophages, e.g., phagocytosis, expression of Fc (fragment, crystallizable region) receptors, and complement receptor (CR)3. The rapid response of microglia to areas of neuronal death and shift in morphology to an amoeboid phagocytic phenotype support early contentions that microglia served as brain macrophages. While T cells actively monitor the brain and can be recruited to sites of injury or infection, they do not remain or become functionally active unless a stimulating antigen is present. In the CNS, infiltrating and resident myeloid cells express MHC class II (Carson et al. 2006; Siffrin et al. 2007), while neurons, astrocytes, and oligodendrocytes are not players in antigen-mediated activation or retention of CD4+ T cells. Similarities between peripheral and central macrophages have led to assumptions that the innate immune system in the CNS equates to the parallel peripheral system and that expression of various factors such as MHC II or cytokines by resident cells of the brain represents neuroinflammation that can occur in the absence of peripheral immune cells. However, this independent capability of CNS immune cells comes under question with regard to whether the cellular response of microglia demonstrates neuroinflammation or simply the diverse nature of the cell (Graber et al. 2011).

Microglia are distinct from other tissue macrophages by their relatively quiescent phenotype and tight regulation by the microenvironment. Microglia display a small amount of perinuclear cytoplasm; small, dense, and heterochromatic nucleus; and several long processes with multiple branches covered with very fine protrusions. They are located outside of the vascular basement membrane; however, the cytoplasmic processes intermingle with astrocytic foot processes in the perivascular glia limitans (Lassmann et al. 1991). Perivascular microglia are limited along vascular walls and distinct from perivascular cells (Graeber et al. 1990). Microglia comprise approximately 15–20 % of the total number of cells in the mature brain. Occupying defined spatial territories, they actively survey the surrounding parenchyma via dynamic movement of processes, while the soma remains static, allowing each cell to sample its environment over the course of a few hours (Davalos et al. 2005; Nimmerjahn et al. 2005). The physical extent of this surveillance depends upon distribution, individual cell ramification, regional location, and rate of process remodeling.

### 2 Distribution of Microglia

Differences in microglia localization and morphology (Fig. 1) are suggested to be associated with functional differences and the phenotypic heterogeneity, including receptor expression patterns (Olah et al. 2011). Early evaluations by Lawson et al. (1990) showed a relatively uniform distribution of microglia in the adult rodent brain, with the exception of a higher relative number of microglia with similar phenotype in the dentate gyrus of the hippocampus, the substantia nigra (SN), and portions of the basal ganglia and the highest number of microglia in the olfactory telencephalon. Microglia modify their morphology and expression of cell-surface antigens depending upon their environment (Kaur and Ling 1991). Ramified microglia in gray matter show processes extending in multiple directions, while in the white matter, cells align cytoplasmic extensions in parallel or at right angles to nerve processes. In regions lacking a blood–brain barrier (BBB) such as circumventricular organs, a less ramified morphology is observed (Perry and Gordon 1987).

### 3 Microglia Network and Response

Microglia are in constant contact with other cells of the brain maintaining interactions with astrocytes, oligodendrocytes, and neurons, as well as brain vasculature. However, the cells do not normally demonstrate a direct intercellular coupling such as through gap junctions (Eugenin et al. 2001); therefore, communication is considered to rely heavily on auto- and paracrine signaling. It is thought that microglia are maintained in a relatively quiescent state due to signals derived from neuronal and astrocyte-derived factors specifically targeted to this regulatory task (Biber et al. 2007; Cardona et al. 2006; Simard et al. 2006). Healthy neurons are able to maintain microglia in an inactive state via secreted and membranebound signals including CD200, CX3CL1 (fractalkine), neurotransmitters, and neurotrophins (Biber et al. 2007; Neumann 2001; Pocock and Kettenmann 2007). In the absence of these signals or in the presence of stimulatory signals, the dynamic surveillance feature allows microglia to rapidly respond to changes in the environment. This is demonstrated in mice deficient of CD200 in which microglia exhibit a less ramified and shorter processes morphological phenotype, increased expression of CD11b and CD45, and elevated production of inflammatory mediators following immune challenge (Hoek et al. 2000).

With stimulation, microglial response patterns are highly varied across morphological changes. Often the initial response of microglia is to monitor and control the extracellular environment. These ever-changing phenotypes represent the unique nature of the insult as well as the ability of the cells to respond in a graded manner to changes in their environment (Graeber and Streit 2010) and to shift within any specific stage of the response (Stout et al. 2005). The phagocytic feature of these cells serves to remove excess cellular debris and aberrant proteins such as apolipoprotein E (ApoE), mutant huntingtin, or amyloid beta 42 (A $\beta_{42}$ ).

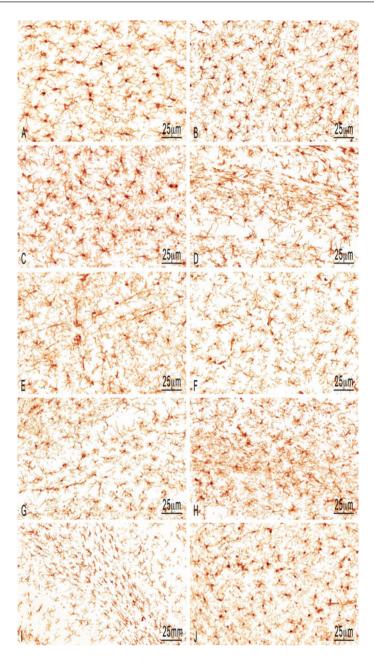


Fig. 1 Representative Iba-1+ microglia in various regions of the rat brain. (a) Frontal cortex, (b) motor cortex, (c) retrosplenial agranular cortex, (d) corpus callosum, (e) striatum, (f) thalamus, (g) hippocampal dentate hilus, (h) hippocampus molecular layer, (i) cerebellum, (j) olfactory bulb. Images from 40 m sections were scanned at  $20 \times$  magnification using an Aperio Scanscope T2 Scanner (Aperio Technologies, Inc., Vista, CA) and viewed using Aperio Imagescope v. 6.25.0.1117

Microglia activities observed in neuroimmunological disorders, such as multiple sclerosis, and their related function to remove myelin debris are a classical representation of the cell's macrophage and antigen-presentation role. This is characterized by prototypical pathology dominated by perivascular mononuclear infiltrates, residual fibrosis of blood vessels, and demvelinating lesions (Barnett et al. 2009; Stadelmann 2011). In an autoimmune disease such as MS, microglia serve in reactivation of infiltrating T cells (Coyle 2011). In contrast, microglial activation observed with a sterile insult does not involve antigen-based interactions with T cells or a predominant infiltration of peripheral immune cells. A strong microglia reaction occurs at the primary site of injured neurons progressing temporally and spatially to recruit responses in areas that are projection sites to and from the area of damage (Graeber and Kreutzberg 1988; Jensen et al. 1999; Sanders and Jones 2006). While anterograde and retrograde degeneration of neuronal projections most likely causes a microglial reaction in the first-order projection areas, other mechanisms, like neuronal ionic balance, likely drive the microglia response in more distant and remote areas. Even within a focal injury site, a graded level of response can be observed. For example, with an ischemic or physical lesion to the brain, the microglia rapidly wall off areas of the CNS from non-CNS tissue and remove dead, damaged, or dysfunctional cells, while in the periphery of the injury, the cells likely perform a survival function.

### 4 Cell-Responsive Interactions

While a number of mechanisms have been proposed for cell-responsive interactions of microglia, many associated with ionic balance are related to the presence of inwardly rectifying potassium channels (Kettenmann et al. 1993) and multiple neurotransmitter receptors (Pocock and Kettenmann 2007). Microglia can be induced to produce several agonists of excitatory amino acid receptors including glutamate, quinolinate, and D-serine (Barger 2004). Several types of glutamate receptors are expressed, including the mGluR (metabotropic glutamate receptor) and AMPA (alpha-amino-3-hydroxy-5-methylisoxazole-4-propionic acid) classes (D'Antoni et al. 2008) and individual receptor subunits of the metabotropic groups I, II, and III (Byrnes et al. 2009; Noda et al. 2000). Activation of mGluR from group I or III inhibits microglia activation; group II evokes mal-activation, and activation of mGluR3 inhibits toxicity of microglia toward oligodendrocytes (Pinteaux-Jones et al. 2008). Recent work demonstrated that activation of Gai-coupled receptors, including group II and group III mGLuRs, suppresses the export of glutamate from activated microglia (McMullan et al. 2012).

A multitude of signals, both physical and secreted factors, posing a potential threat to brain homeostasis are sensed by microglial receptors (Hanisch and Kettenmann 2007). Microglial activation, in addition to being stimulus dependent, is also likely to be a multistep process (Ponomarev et al. 2006). Specific factors released by stressed or damaged neurons can stimulate the production of proinflammatory cytokines by microglia including matrix metalloproteinase-3 (MMP-3),  $\alpha$ -synuclein, neuromelanin, and adenosine triphosphate (ATP). In an in vitro transmigration assay, conditioned media from mesencephalic cultures

treated with MPP+ (1-methyl-4-phenylpyridinium) were sufficient to trigger microglia attraction. This response occurred even at early stages of the neuronal injury, suggesting signaling from the neuron to trigger microglia activation. The rapid response to injury of directed process extension relies on purinergic signaling mediated by ATP and P2 (purinergic) receptors. In addition to local process extension, microglia can sense a chemical gradient of ATP and exhibit chemotaxis to the source. The P2Y12 receptors mediate chemotaxis and the P2Y6 receptors stimulate phagocytosis (Koizumi et al. 2007; Langosch et al. 1994).

Depending upon the stimulus, inflammatory responses by microglia can be initiated by pattern recognition receptors (PRRs) that include Toll-like receptors (TLRs), the receptor for advanced glycation end products (RAGE), and scavenger receptors to detect the aberrant expression of phosphatidylserine on the extracellular surface of dying cells. Microglia detect ligands for CD40, CD91, and the intracellular NOD-like receptors (NLRs) that initiate the signaling process by binding to PAMPs. Ligation of PRRs leads to the activation of signal transduction pathways and regulation of diverse transcriptional and posttranscriptional molecules including members of the nuclear factor kappa B (NF- $\kappa$ B), activator protein 1, and interferon regulator factor families, which modulate proinflammatory target genes encoding cytokines, chemokines, enzymes, and other molecules essential for pathogen elimination (Akira et al. 2006).

TLRs are a major family of PRRs for a diverse set of novel pathogen-associated molecules (Takeuchi and Akira 2001) that, with their related signaling proteins, are expressed in the CNS (Bowman et al. 2003; Kielian et al. 2002) with microglia expressing TLRs 1–9 (Farina et al. 2005). In vivo, TLR2 expression is exclusively in microglia activated with cerebral ischemia (Lehnardt et al. 2007) or entorhinal cortex lesion (Babcock et al. 2006). Microglia derived from human white matter and in primary rat cell cultures express TLR4 (Lehnardt et al. 2003; Visintin et al. 2001) that recognizes the fragment of gram-negative bacteria and LPS and can initiate innate immune responses to infection in mammals. A large proportion of the data suggests that TLR signaling mediates beneficial effects essential for pathogen elimination, but additional data suggests that TLR-induced microglia activation and release of proinflammatory molecules can contribute to neurotoxicity. Based upon studies examining the various TLRs, it has been suggested that activation of innate immune responses in the brain is tailored according to the cell type and environmental signal. As an example, TLR3 signaling induces a strong proinflammatory response in microglia as characterized by the secretion of IL-12, TNF- $\alpha$ , IL-6, CXCL-10, and interferon (IFN)-β. TLR2-mediated responses are primarily associated with secretion of IL-6 and IL-1 $\beta$  (Jack et al. 2005).

More recently, PRRs have been found to respond to factors released from necrotic cells and by molecules that may be secondary to a pathogenic process and may facilitate neuronal damage. TLRs, especially TLR2 and TLR4, are capable of sensing damage induced by ischemia and, as such, boost the proinflammatory response such that the infarct size is increased (Caso et al. 2008; Kilic et al. 2008; Lehnardt et al. 2007). There are several host-derived ligands for TLR. One of these, heat shock protein (HSP)60, is released from dying CNS cells and, upon binding to

microglia, TLR4- and MyD88-dependent secretion of NO is induced (Lehnardt et al. 2008). Similarly, necrotic neurons have been shown to activate microglia in an MyD88-dependent manner with the subsequent proinflammatory response leading to an increased neurotoxic activity through the induction of the glutamate enzyme, glutaminase (Pais et al. 2008). While TLR activation can contribute to neurotoxicity during CNS infection, there is evidence that TLR signaling can mediate beneficial effects (Glezer et al. 2006). Microglia appear to be the major initial sensors of danger or stranger signals recognized by TLR4, and they secrete inflammatory mediators such as TNF- $\alpha$  and IL-1 $\beta$ . These cytokines can then act on astrocytes to induce a secondary inflammatory or growth factor repair response (Saijo et al. 2009). With an acute injury, the release of TNF from microglia may serve to counter any secondary injury (Nawashiro et al. 1997), while activation of microglia to remove cellular debris following acute damage may serve to prevent subsequent tissue inflammation (Glezer et al. 2007; McPherson et al. 2011). TLR4 is also used by microglia and astrocytes to detect AB (Landreth and Reed-Geaghan 2009), leading to activation of signal-dependent transcription factors for downstream inflammatory response genes and clearance of aberrant proteins (Reed-Geaghan et al. 2009; Walter et al. 2007). Supporting data suggests that stimulation of the system with TLR ligand encourages the appropriate phenotype of resident or invading monocytes to promote the clearance of A $\beta$  (Yong and Rivest 2009).

RAGE, a cell-surface receptor belonging to the immunoglobulin superfamily (Neeper et al. 1992; Schmidt et al. 1992), is present on the surface of microglia, astrocytes, vascular endothelial cells, and neurons. Activation occurs with production of advanced glycation end products in prooxidant and inflammatory environments. RAGE contributes in the clearance of A $\beta$  and is involved in apolipoprotein E (ApoE)-mediated cellular processing and signaling. It recognizes other ligands, including serum amyloid A, S100 protein, and high-mobility group box 1. NLRs are soluble, cytoplasmic PRRs that act as sensors of cellular damage. In Alzheimer's disease (AD), A $\beta$  oligomers and fibrils induce lysosomal damage, triggering the NOD-like receptor, NALP3, in microglia (Halle et al. 2008). NALPs activate downstream signaling proteins, such as apoptosis-associated speck-like protein containing a caspase recruitment domain. This induces apoptosis, but it also contributes to the maturation of proinflammatory mediators like IL-1 $\beta$  and IL-18. In addition to peptide fragments, a decrease in cellular potassium concentration can activate NALP1 in neurons, leading to a similar activation of ASC, apoptosis, and IL-1 $\beta$  and IL-18 maturation.

Additional receptors that recognize apoptotic cellular material, such as phosphatidylserine on the inner membrane leaflet, are important for phagocyte clearance processes and may stimulate an anti-inflammatory response (Ravichandran 2003). Recent identification of such receptors includes T-cell immunoglobulin- and mucindomain-containing molecule-4 (TIM4) (Miyanishi et al. 2007), the metabotropic P2Y6 receptor that recognizes the nucleotide UDP released from injured neurons (Koizumi et al. 2007), and the triggering receptor expressed on myeloid cells-2 (TREM-2; Schmid et al. 2002) to facilitate debris clearance in the absence of inflammation (Takahashi et al. 2005). A critical role for TREM-2 signaling has been demonstrated in polycystic lipomembranous osteodysplasia with sclerosing leukoencephalopathy, or Nasu-Hakola disease, with early-onset dementia possibly due to the inability of microglia to clear tissue debris via TREM-2 signaling (Neumann and Takahashi 2007). The differential expression of these receptors in response to the intensity or stage of tissue injury, type of injurious stimuli, and presence of other soluble signals can exert significant control over the potency of the microglial response.

### 5 Microglia Cytokine Expression and Regulation

Microglia are a source of a number of cytokines such as IL-1, IL-6, and TNF and are responsive to growth factors such as granulocyte-macrophage-CSF and CSF-1. Cytokines can induce or modulate a broad spectrum of cellular responses including cell adhesion, migration, survival, differentiation, replication, secretory function, and cell death. In the same cell, a cytokine can either induce death or promote survival or proliferation depending upon the function of the context in which it acts. Cell death can be actively triggered by a cytokine as a physiologically necessary role during development, differentiation, and in host defense responses to injury and subsequent repair. However, the same cytokine can also provide critical survival factors, and thus, its absence can result in cell death at a specific point in the cell cycle.

A specific phenotype or function has yet to be directly linked to specific altered or activated microglia morphology. A specific classification of microglia phenotypes has been proposed for microglia cells based upon peripheral myeloid cell responses and cytokine release (Martinez et al. 2008; Michelucci et al. 2009; Mosser and Edwards 2008). For example, proinflammatory (Th1, Th17) CD4+ T cells promote myeloid cells toward a classical activation state (M1 state) characterized by high expression levels of proinflammatory cytokines (IL-1 $\beta$ , IL-12, IL-23, IL-6, TNF- $\alpha$ ) (Colton and Wilcock 2010; Gordon and Martinez 2010; Graeber and Streit 2010). In contrast, Th2 CD4 + T cells promote myeloid cells toward alternative activation states (M2a and M2b) characterized by expression of macrophage mannose receptor (CD206), YM-1, stabilin-1, and arginase I. Under this alternative activation, cells are functionally associated with enhanced phagocytosis, tissue repair, and parasite elimination. A regulatory activation state (M2c) is promoted by IL-10 and TGF $\beta$ characterized by expression of molecules related to tissue repair and immunosuppression. Each state may have its own particular phenotype with likely transitional phenotypes between wound healing classical and regulatory macrophages (Mosser and Edwards 2008). While this stage distinction has been established for peripheral cells, whether this will translate to the CNS has yet to be clearly established, as neither a temporal nor a spatial transition from an M1 to an M2 phenotype has been clearly described in models of neuroinflammation or neurodegeneration (Ajami et al. 2007; Colton and Wilcock 2010; Hume et al. 1983). Rather, as a general pattern, data suggest heterogenous populations of M1 and M2 microglia throughout an injury response or neurodegenerative disease process.

To understand the impact of a cytokine response, the basal level and regulatory mechanisms need to be considered. Under normal homeostatic conditions, the majority of cytokines are either not expressed or are expressed at low levels. In the early phase, host defense responses to injury are triggered by microbial antigens, extracellular matrix antigens, or by activation products or preformed components of humoral defense systems. This process consists of cell recruitment and induction of cytokines that contribute to activation of proinflammatory, immunostimulatory, and catabolic responses. Cytokine mRNA transcripts have a short half-life of 10–30 min; thus, cytokines are rapidly translated and secreted into the extracellular space, producing a burst of cytokine release. Some cytokines are secreted upon complete intracellular processing, while others are stored intracellularly, and additional stimuli are required to trigger secretion. For example, transcription without translation can be observed for IL-1 following such things as adherence of blood monocytes to surfaces or exposure to calcium ionophore. In these cases, steady-state mRNA levels for IL-1 $\beta$  are comparable with those following induction by endotoxin; however, there is no protein translation. Alternatively, the system may effectively regulate the downstream effects of IL-1 by the induction of IL-1 receptor antagonist to bind the protein and prevent receptor activation.

Most cytokines are biologically active when secreted; however, some, such as TNF- $\alpha$  and transforming growth factor (TGF) $\beta$ , are secreted in a biologically inactive or latent form and require further extracellular processing by proteases. Cytokines regulate the expression of cytokine receptors involved in the modulation of activity. For IL-1 and TNF- $\alpha$ , these effects are mediated primarily by IL-1 type I receptor and TNF receptor TNFR1 (p55) or TNFR2 (p75), respectively. IL-6 and other members of this family, oncostatin M, leukemia inhibitory factor, IL-11, and ciliary neurotrophic factor, are induced by IL-1 and TNF and protect against the proinflammatory and catabolic effects. As one primary example, microglial functions related to an innate immune response are associated with TNF signaling. TNF can induce apoptosis via activation of receptors containing an intracellular death domain such as TNFR1 and CD95 (APO-1/Fas) with their corresponding death ligands, TNF, and the structurally related type II transmembrane protein, FasL. Release of TNF- $\alpha$  and FasL shedding by microglia are implicated in neurotoxicity. Signaling by TNF is dependent upon constitutive and induced target cell expression of TNFRs (Tartaglia and Goeddel 1992). Membrane receptor mechanisms of apoptosis are implicated in neuronal death, while soluble TNFR1 can offer protection by binding TNF and preventing receptor signaling. TNFR1 activation has been proposed as a molecular mechanism for rapid apoptosis of injured or sick neurons through a caspase 3-mediated pathway (Yang et al. 2002). Recent studies examining cell death and survival following an ischemic insult or induced by a systemic injection of the organometal, trimethyltin, suggest that the level of TNF- $\alpha$  produced by microglia and the neuronal expression pattern of TNF receptors serve as critical determining factors for neuronal death (Harry et al. 2008).

### 6 Release of Factors Associated with Oxidative Stress

In vitro, microglia are capable of releasing several potentially cytotoxic substances, such as free oxygen intermediates, NO, proteases, arachidonic acid derivates,

excitatory amino acids, quinolinic acid, prostaglandins, cytokines, and other factors that, if occurring in vivo, may contribute to the exacerbation of neurons already compromised by disease or injury. In parallel with the production of proinflammatory cytokines, microglial neuroinflammation is commonly associated with the production of ROS and NO-dependent reactive nitrogen species (RNS). In excess concentrations, NO forms peroxynitrite and, through nitrosylation of cell signaling messengers, functions in immune regulation (Guix et al. 2005). ROS are oxygen-containing molecules that react with and oxidize vulnerable cellular constituents, including proteins, nucleic acids, and lipids. In neural tissues, oxidative stress can result in disrupted signaling processes and ion homeostasis and is held accountable for events ranging from protein mis-folding to death of newly generated neurons (Lipton et al. 2007; Taupin 2010). Superoxide, in particular, is implicated in microglial activation, cellular redox imbalance, and associated neurodegeneration. The translation of these processes to humans is hindered by the fact that levels induced by activating stimuli significantly differ across species and are relatively low in humans (Colton et al. 1996).

Endogenously produced ROS and RNS are essential components of brain homeostatic processes. ROS are commonly released by microglia to eliminate pathogens and elicit the controlled destruction of neuronal debris. In general, superoxide production by myeloid cells is mediated by the phagocytic NADPH oxidase (NOX2), consisting of membrane-bound (gp91 and p22) and cytosolic (p47, p67, and p40) subunits, as well as a requirement for the GTPase, Rac, for full activity (Lambeth 2004; Sumimoto et al. 2004). The microglial respiratory burst in response to certain tissue-disrupting stimuli involves Ca<sup>2+</sup> and K<sup>+</sup> channels and NADPH-dependent signaling to induce the release of superoxide (Colton et al. 1994; Khanna et al. 2001). Opening of cell-surface ion channels on microglia specific for induction of ROS as opposed to NO, including P2X7, Kv1.3, TRPV1, and KCa3.1, causes an "acute phase" of activation, often involving protein kinase C (PKC)-dependent signaling to NADPH oxidase and downstream NF- $\kappa$ B prior to superoxide release and gene induction (Fordyce et al. 2005; Infanger et al. 2006; Parvathenani et al. 2003; Schilling and Eder 2010).

NOX2 can be activated in mononuclear cells by TNF- $\alpha$ , IFN- $\gamma$ , IL-1 $\beta$ , prion protein, ATP, and fibrillar A $\beta$ . In addition, although phagocytosis can cause activation of NOX2, microglia do not uniformly undergo a respiratory burst when they initiate a phagocytic action (Savill et al. 2003). Notably, activation of NOX2 alone is not always sufficient to cause neurotoxicity (Mander and Brown 2005), suggesting that microglial superoxide production may have to occur in connection with other stressors to induce neurodegeneration. Some of these other factors that may contribute to neurodegeneration include free radicals (i.e., hypochlorous acid and reactive nitrites) produced via myeloperoxidase activity in astrocytes or peripheral leukocytes (Choi et al. 2003; Chung et al. 2010), and ROS-stimulated microglial proliferation and cytokine production (Jekabsone et al. 2006; Mander et al. 2006). The work of Barger et al. (2007) suggested that glutathione (GSH) depletion and oxidative stress resulting from the NADPH-dependent respiratory burst induce glutamate release from microglia, resulting in neuronal loss. Induction of NADPH oxidases has been shown to be involved in the neurotoxic response in various Parkinson's disease (PD)-type models, including LPS, 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP), rotenone, and paraquat, with a possible requirement for protein kinase C (PKC) delta phosphorylation of p67 and p47 (Miller et al. 2009; Qin et al. 2004; Tieu et al. 2003).

A key event in a ROS stress response is the induction of heme oxygenase-1 (HO-1) in microglia and other cells that can help protect against subsequent insults (Lee and Suk 2007). Although increased HO-1 activity and CO production can have neurotoxic effects in response to select insults, the regulated activation of this system in microglia is hypothesized to be involved in resolving neuroinflammation and preventing accessory tissue damage (Syapin 2008). While NO-mediated stress is implicated in the progression of multiple brain insults, including AD, PD, and stroke (Guix et al. 2005), it can also induce neuroprotective signaling events such as inactivation of caspases, modulate release of neurotransmitters (e.g., dopamine, acetylcholine, GABA, and glutamate), and regulate synaptic plasticity by facilitating the induction of long-term potentiation via soluble guanylyl cyclase (Bon and Garthwaite 2003; Calabrese et al. 2007). Inducible NOS (iNOS; NOS2) present in microglia is Ca<sup>2+</sup> independent and can be upregulated in response to TNF- $\alpha$  and IFN- $\gamma$  or downregulated in response to TGF- $\beta$  and IL-1 $\beta$  (Tichauer et al. 2007). Microglial NO can be cytoprotective and neuroprotective against select insults, such as ischemia. At high levels, iNOS can be cytotoxic to oligodendrocytes and can lead to excitotoxicity and/or synapse strengthening and influence BBB permeability. Thus, the induction of an oxidative stress response in microglia may likely underlie much of the neurodestructive effects attributed to microglia activation.

### 7 Microglia Aging and Senescence

The progressive, age-related change in MHC II antigen expression by microglia led to the assumption that microglial activation occurs with aging. The expression pattern was reported as similar to what occurs in experimental animal models of injury (Letiembre et al. 2007; Perry et al. 2002). Systemic challenge of aged mice with LPS suggested a primed phenotype for the microglia based upon the elevated basal level of cytokine synthesis as compared to young animals (Dilger and Johnson 2008). Further investigation showed morphological changes of microglia cytoplasmic structure in the aged human brain reflective of dystrophy and senescence but no evidence of priming (Sheng et al. 1998). Thus, while the cells change as a function of age, the morphological phenotype is suggestive of a decreased ability to mount a normal host-resistance response rather than a transformation to an activated cell. It has been suggested that age-related neurodegeneration might be due to not only a loss of neuroprotective properties but also the actual loss of microglia (Streit 2006; Streit and Xue 2012). One of the more prominent features of microglia is their function to clear aberrant proteins and debris from the CNS. Thus, the loss of efficiency in this function would result in a more toxic microenvironment.

A recent hypothesis put forth to explain phenotypic changes in aging microglia suggests that microglial senescence is triggered by an intracellular oxidative stress response due to enhanced intracellular accumulation of iron (Streit 2006). Under this hypothesis, a prolonged oxidative stress response would lead to senescence rather than activation. Additional support for microglial replicative senescence in normal and pathological aging comes from the demonstration of telomere shortening and decreased telomerase activity with aging. Whether this occurs with agerelated diseases remains in question; however, data suggest that microglia from AD brains have shorter telomerase, as compared to control brains (Streit 2006). Using A clearance as a phagocytic function of microglia that may be compromised with aging, a number of studies have examined a possible cellular process that could contribute to plaque accumulation in aging brains. Microglia cultured from aged mice showed a diminished ability to internalize A $\beta$  peptide. CD47-dependent ability to phagocytize AB fibrils seen in postnatal day 0 microglia was lost in microglia isolated from 6-month-old mice and appeared to manifest between the ages of 2 and 6 months. This effect was not due to a generalized loss of phagocytic capability as no age-related difference was observed in clearance of bacteria but rather appeared specific for A $\beta$ . In mouse models of AD, the increased plaque deposition that can occur as early as 6 months of age may be related to the loss of selective phagocytic function of microglia (Westerman et al. 2002). It has also been reported that between 6 and 18 months of age, microglia show an increase in mRNA levels for the alternative activation marker, YM-1. By comparison in a transgenic AD mouse line expressing human mutant amyloid precursor protein (APP) and presenilin 1 (PS1), significantly higher levels of YM-1 mRNA were demonstrated at 6 months of age (Jimenez et al. 2008). In contrast to previous reports suggesting that mRNA levels for proinflammatory cytokines increased with age, AD transgenic but not wild-type microglia displayed decreased Aβ phagocytic ability and increased mRNA levels of classic activation markers such as TNF-a at 18 months in non-plaque areas of the brain. One hypothesis put forth is that interactions of A $\beta$  peptide with other inflammatory stimuli in the diseased brain drive the cells to an early and maintained alternative activation state that shifts the ability of the cells to respond in a normal manner (Floden and Combs 2011). In support of an age-related decrease in responsiveness of microglia, ApoE4 transgenic mice displayed a deficit in the resident microglia host-resistance and proinflammatory cytokine response to injury as a function of age (Harry et al. 2000). The deficit was not observed at 2 months of age but only became apparent at 8 months of age.

### 8 Acute Injury

Several studies demonstrate that, with acute injury, microglia can be destructive or protective (Butovsky et al. 2005; Nelson et al. 2002). With an acute physical disruption of the brain, the cellular response of the brain macrophage is primarily limited to the immediate focal site and the surrounding parenchyma. With a slightly

deeper penetration to include the corpus callosum, responses along the myelinated tracts can be observed. The nature of a puncture wound injury involves recruitment of blood-borne cells to the reaction including leukocytes, granulocytes, and monocytes. Within the lesion itself, round hematogenous macrophages are primarily found. At the edge of the lesion, process-bearing, hypertrophic, resident microglia are evident. This initial response occurs within the first 12 h or so, peaks between days 2 and 3, and by day 5, hypertrophied microglia express MHC class II and the CD4 antigen. This response then declines over the next 5 days. With the injection of IFN- $\gamma$ /LPS, the response is slightly modified given the more active recruitment of peripheral immune cells. Interactions with CD4+ T cells with influence microglial phenotype and effects on neurons with fewer neurotoxic factors produced and an increase in anti-inflammatory and growth mediators and a reverse of glutamate transporters resulting in a buffering rather than a release of glutamine (Shaked et al. 2004, 2005). Thus, acute injury models offer critical information on the interaction between peripheral and central immune cells for consideration in more delayed neuronal death or progressive neurodegenerative models.

### 9 Progressive Degenerative Disease States

While the presence of reactive and activated microglia has been observed in numerous animal models of human neurodegenerative disease and in patients, the overall evidence suggests that, in the context of a pathogen-free environment, a relatively benign innate immune response is evoked. However, how, and if, this environment can be altered or stressed in a manner to change this balance remains a critical question. In murine prion disease models, particular prion agents can induce pathology in specific regions of the brain. The pathology is often characterized by protease-resistant protein deposition, spreading through the brain, accompanied by an increase in morphologically activated microglia with hypertrophied cell bodies. In the ME7 model in C57BL mice, at the time of first detectable synaptic loss, microglia in the hippocampus show an activated morphology and elevated expression of antigens such as CR3, ERM1, and CD68 (Betmouni et al. 1996). With further activation and during the progression to terminal stages of the disease, an increase is observed in TGF<sup>β</sup>, cyclooxygenase (COX)-2, and its product PGE2 and the chemokine CCL2/MCP-1 (Perry et al. 2002), which often function in an anti-inflammatory manner. While the expression of TGF $\beta$  and PGE2 represents macrophages that have phagocytosed apoptotic often cells (Savill et al. 2002), there is no evidence of neuronal apoptosis in these early stages. It is possible that the activation occurring with synapse loss reflects a microglia phenotype as a consequence of clearing degenerating synapses; however, in this model, it is the dendritic spine that envelops the degenerating synapse and not the microglia (Siskova et al. 2009).

A similar benign role has been suggested in models of AD. In postmortem brain tissue of AD patients, an atypical inflammation occurs with both activation of the resident microglia and infiltrating monocytes surrounding dense-core amyloid- $\beta$ 1-40/42 plaques showing increased expression of cell-surface antigens (D'Andrea et al. 2004; McGeer and McGeer 2003; Simard et al. 2006). In transgenic mice expressing human APP, amyloid plaque formation and amyloidassociated neuritic dystrophy occur independent of microglia presence (Grathwohl et al. 2009). In many of these animal models, microglia are a major component of the dense-cored amyloid plaques but not diffuse plaques, and it has been hypothesized that in the absence of effective phagocytosis, microglia potentiate the toxicity of A $\beta$  peptide (Streit et al. 2005). Current data support the idea that inflammatory mediators do not promote AB accumulation but limit AB deposition (Chakrabarty et al. 2010). While the phagocytic activation serves in a beneficial manner to eliminate aggregated and toxic proteins from the brain, it has been proposed that an extended activation or dysregulation of this system can lead to an adverse environment and toxicity to surrounding cells. A recent study using the Tg-SwDI mice as a model for AD demonstrated that activated microglia localized around microvascular amyloid deposits increased synthesis of native complement proteins C1q, C3, and C4 (Fan et al. 2007). In transgenic mouse models of AD, less neuropathology is demonstrated in the absence of C1q; however, inhibition of complement activation diminishes microglial activity, increases amyloid load, and enhances neuronal degeneration (Fonseca et al. 2004; Wyss-Coray et al. 2002).

In PD, dopaminergic and non-dopaminergic neurons in the nigrostriatal pathway are compromised, contributing to the motor-related symptoms. A number of mechanisms have been proposed for the death of dopamine-containing neurons in the SN pars compacta including mitochondrial dysfunction, oxidative stress, and impairment of protein degradation. The detection of elevated levels of proinflammatory cytokines and evidence of oxidative stress-mediated damage in postmortem PD brains gave rise to the hypothesis of a microglia involvement. However, observations of the presence of cytotoxic T cells in the SN of a patient with PD and an increased density in insulin-like growth factor  $\gamma$ + cells in the brains of PD patients suggest a role for the cellular arm of the adaptive immune system. A critical role for T cells was demonstrated in two different strains of immunodeficient mice lacking mature T lymphocytes (Tcrb-/- and Ragl-/- mice) by a reduction in MPTPinduced dopaminergic cell death. Further work demonstrated CD4+ T cellmediated cytotoxicity of MPTP (Brochard et al. 2009) possibly via microglia activation and release of free radicals and proinflammatory cytokines. However, the recent description of an IL-17 secreting Th17 lymphocyte suggests an alternative cellular source for proinflammatory cytokines. One could envision that with progressive changes in the neuron, a microglia response to  $\alpha$ -synuclein occurs, recruiting CD4+ T cells, leading to an induction of proinflammatory cytokines and NO secretion by microglia alternatively or in conjunction with the infiltration of TH17 lymphocytes. When a progressive model of PD was examined with reference to the adverse impact of microglial activation and inflammation on dopaminergic neurons, Wright and colleagues (2008) demonstrated in vivo that while the immunosuppressant drug, tacrolimus (FK506), reduced the initial stage of neuronal death in the SN, it did not alter the morphological response of microglia, suggesting a lack of direct correlation between the two cellular responses.

### 10 Conclusion

The fact that microglial activation correlates with neurodegenerative disease progression has led to the hypothesis of a causal link; however, the rapid nature of microglia response to neuronal injury, both physical and chemical signals, and death makes it difficult to determine an initiating role for microglia. There is a critical need to gain a better understanding of the plethora of responses representing the dichotomy of microglial reactivity for beneficial versus destructive effects. Identifying what can trigger a microglia response and identifying characteristics of the various types of responses will further advance the ability to interpret the role being played by microglia under normal and aberrant brain conditions.

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# 6-Hydroxydopamine Lesioning of Dopamine Neurons in Neonatal and Adult Rats Induces Age-Dependent Consequences

# Sophia T. Papadeas and George R. Breese

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S.T. Papadeas (⊠) Chapel Hill, NC, USA

e-mail: spapade129@gmail.com

G.R. Breese

Department of Psychiatry and Pharmacology, Bowles Center for Alcohol Studies, UNC School of Medicine, Chapel Hill, NC, USA

e-mail: george\_breese@med.unc.edu

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#### Abstract

Over 45 years ago, it was discovered that the compound 6-hydroxydopamine (6-OHDA) destroys catecholamine (CA)-containing neurons when injected directly into the brain. Refinements in the use of 6-OHDA led to the development of dopamine (DA)-denervated rats to model Parkinson's disease (PD) and Lesch-Nyhan syndrome (LNS), in which loss of DA-containing neurons is a prominent feature. Differing functional characteristics were eventually discovered in DA-denervated rats depending on the age at which the DA was reduced. With 6-OHDA lesioning of CA neural systems in the brain, key discoveries were made regarding how these systems control specific physiological functions and behavioral responses. Important neurobiological concepts emerged as well, including receptor supersensitivity, D<sub>1</sub>/D<sub>2</sub>-DA receptor interaction, and D<sub>1</sub>-DA priming of supersensitivity, including neural mechanisms likely relevant to clinical symptoms modeled by the 6-OHDA treatment. In this chapter, behavioral, neurochemical, structural, and pharmacological studies carried out in adult and neonatal 6-OHDA-treated rats are reviewed, and findings are compared and contrasted. To date, age-dependent lesioning of DA neurons in rats with 6-OHDA continues to play an important role in the development and validation of new therapies for clinical disorders with brain DA reduced.

#### Keywords

Adult 6-OHDA • DA-receptor interactions • Lesch-Nyhan disease • Neonate-6OHDA • Parkinson's disease

### 1 Introduction

The neurotoxicant 6-OHDA was the first agent to produce long-term depletion and destruction of norepinephrine (NE) in peripheral sympathetic neurons without affecting central CA content (Porter et al. 1963; Laverty et al. 1965; Thoenen and Tranzer 1968). Subsequently, the toxin was found to produce a selective destruction of CA-containing neurons when administered directly into the brain (Ungerstedt 1968; Bloom et al. 1969; Breese and Traylor 1970; Uretsky and Iversen 1970).

Over the last 4 decades, administration of 6-OHDA into the brain has been used effectively in both adult (Ungerstedt 1968; Breese and Traylor 1970; Breese et al. 1974) and neonatal (Breese and Traylor 1972; Smith et al. 1973) animals to investigate age-dependent behavioral, physiological, and biochemical effects of midbrain CAergic neuronal loss in disease symptoms and for identification of drug action. From these efforts, DA and NE neurons were found to be essential for the action of psychostimulant drugs, control of movement, and behavioral processes such as emotion, attention, reward, body temperature, and stress responses. Additionally, the basic information collected with 6-OHDA on brain function allowed for defining the role of these CAs in the pathogenesis and/or symptoms of PD, Lesch-Nyhan syndrome (LNS), and a number of other neuropsychiatric disorders, including attention deficit hyperactivity disorder (ADHD), schizophrenia, and drug addiction.

In the present chapter, an extensive overview of 6-OHDA lesioning in rats is provided, highlighting the distinctions between adult and neonatal CA-depleted animals. First, the history, scientific basis, and application of 6-OHDA lesioning techniques to selectively destroy nigrostriatal DA and/or NE neurons of the adult and neonatal rats are discussed. Next, important contributions from previous investigations regarding the behavioral and compensatory neural responses to agedependent 6-OHDA lesioning of these neurons are reviewed. Focus is placed on characteristics unique to rats lesioned with 6-OHDA to destroy DA neurons during the neonatal period, particularly with regard to the differential sensitivity of these rats to various agonists and antagonists of DA receptors and to changes in other transmitter receptors in comparison to adult DA-depleted animals. Finally, potential mechanisms associated with the neural and behavioral plasticity in 6-OHDAlesioned rats are presented, and the usefulness of these animals to model symptoms and features of specific central disorders is emphasized.

### 2 The 6-OHDA-Lesioned Rat Model

### 2.1 Mechanism of Action of 6-OHDA

6-OHDA does not pass the blood-brain barrier, so it must be directly injected into the brain to destroy central CA-containing neurons (Schober 2004; Bove et al. 2005). As an analog of DA and NE neurons, the toxin is readily taken up by CA neurons through the high-affinity DA and NE transporters, DAT and NET, respectively (Jonsson and Sachs 1970; Ljungdahl et al. 1971; Saner and Thoenen 1971). Once inside the neuron, the 6-OHDA initiates degeneration of CA neurons likely through a combination of oxidative stress and mitochondrial respiratory dysfunction (Schober 2004; Bove et al. 2005). The 6-OHDA-induced destruction of DA or NE nerve endings, and possibly cell bodies (depending on dose), results in the denervation of DA- and/or NE-containing brain neurons, which ultimately induces development of DA and/or NE receptor supersensitivity and other adaptations to compensate for this neural loss.

### 2.2 Microinjections of 6-OHDA into the Brain to Lesion Nigrostriatal DA Neurons in Adult Rats

In 1968, Ungerstedt showed that injection of 6-OHDA into the pars compacta region of the SN (SNpc) of adult rats produced anterograde degeneration of the nigrostriatal DA system, thereby generating the first animal model of PD (Ungerstedt 1968). In subsequent studies to disrupt DA function, 6-OHDA was injected either into the medial forebrain bundle (MFB), through which the nigrostriatal pathway ascends, or into the striatum, where axon terminals of SNpc neurons are located (Deumens et al. 2002). With high doses of 6-OHDA (>6  $\mu$ g) injected bilaterally into either the SNpc or MFB of adult rats, nigral DAergic cell death begins within 12 h of injection, followed by a near-complete lesioning of nigrostriatal DA neurons (>90 %) within 2–3 days (Ungerstedt 1968; Jeon et al. 1995; Zuch et al. 2000) – a loss of neurons which mirrors the pattern of DA loss seen in postmortem brains of PD patients. After injection into the SNpc, neighboring DA neurons in the ventral tegmental area (VTA), which give rise to the mesolimbic and mesocortical pathways, are somewhat compromised by 30–40 % as well (Perese et al. 1989; Carman et al. 1991).

Bilateral injections of 6-OHDA into the SNpc or MFB are usually avoided due to a profound acute aphagia and adipsia that follows – an approach which can result in mortality if rats are not tube-fed to maintain nutrition (Ungerstedt 1968, 1971a). Rather than bilateral site injections of 6-OHDA, administration is typically performed unilaterally, with the opposite, or contralateral, side serving as the control. This approach is particularly suitable for evaluating turning behavior to a drug challenge (Ungerstedt 1971d). However, following unilateral lesioning of the nigrostriatal DA pathway, it should be noted that the rat initially exhibits a severe body circling posture to one side for a few days, after which time this deficit nearly subsides – a recovery that obviously depends upon a neural adaptation to allow this functional correction. Consequently, for defining the basis of this adaptation in these unilateral 6-OHDA-lesioned rats, the inclusion of an unlesioned rat would be wise. Robinson and colleagues have provided extensive data documenting the need for appropriate controls to understand the basis of the initial adaptation to unilateral 6-OHDA lesioning in adult rats (Robinson 1991).

Another approach to investigate the consequences of lesioning DA neurons was to inject 6-OHDA ( $12-25 \mu g$ ) into the striatum to produce a more selective and slowly developing partial lesion of nigrostriatal DA neurons (50-70 %) in this site to mimic earlier stages of PD. Though damage to the striatal DAergic terminals is acute (within 24 h of injection) with this method, degeneration of the parent nigral cell bodies begins at 5–7 days postinjection and progresses for at least 4 months after the insult (Sauer and Oertel 1994; Przedborski et al. 1995; Blandini et al. 2007; Berger et al. 1991). This protracted time course of degeneration has provided an in vivo experimental platform for studying neuroprotective strategies aimed at slowing or halting the degeneration of DA neurons in PD (Deumens et al. 2002). Since central administration of 6-OHDA can also reduce brain NE, direct injections of 6-OHDA into the locus coeruleus, the region of the

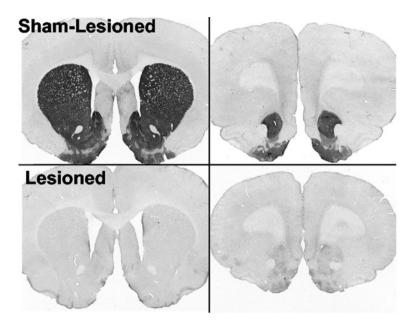
brainstem where NE neurons originate (Ungerstedt 1971c), have also been performed (Ungerstedt 1971e). This strategy of lesioning NE neurons has allowed for the determination of what functional role these particular neurons play in brain function.

### 2.3 Intracisternal Injection of 6-OHDA to Lesion CA-Containing Neurons in Adult Rats

Subsequent to the initial work by Ungerstedt (1968), Breese and Traylor administered 6-OHDA into the cisterna magna of adult rats to produce lesioning of CA-containing neurons (Breese and Traylor 1970). This initial use of intracisternal (i.c.) administration of 6-OHDA to lesion CAergic neurons in the brain was then followed by others (Bruno et al. 1984, 1987; Lytle et al. 1972; Smith et al. 1973). Because initial unpublished efforts demonstrated that 6-OHDA i.c. utilization did not produce the profound reduction of DA observed with injection of 6-OHDA into the SNpc (Ungerstedt 1968), Breese and Traylor pretreated rats with the monoamine oxidase (MAO) inhibitor, pargyline, to produce a comparable reduction of brain DA as seen when 6-OHDA was microinjected into the MFB (Breese and Traylor 1970). Like pargyline, several MAO inhibitors were subsequently demonstrated to also enhance the destruction of DA-containing neurons (Breese 1975). Additionally, the degree of destruction of DA neurons was found to be further enhanced if an additional dose of 6-OHDA was administered 7-10 days after the initial pargyline/6-OHDA exposure (Breese and Traylor 1971; Breese et al. 1973, 1974). Because the 6-OHDA also shows high affinity for the NE transporter, NET (Luthman et al. 1989b), systemic injection of the NET inhibitor, designamine, administered prior to 6-OHDA ensures increased selectivity of the toxin for DA neurons (Evetts and Iversen 1970; Breese and Traylor 1971). Owing to the high affinity of 6-OHDA for NET (Porter et al. 1963), Breese and Traylor found that small doses of 6-OHDA administered i.c. to adult rats reduced brain NE with little effect on DA content in the brain (Breese and Traylor 1971). These approaches allowed evaluating the distinct functions related to either DA or NE in the brain (Breese et al. 1974; Ungerstedt et al. 1974), a strategy which provided a critical understanding of how the loss of CA neurons differentially affected brain functions.

### 2.4 Intracisternal Injection of 6-OHDA to Lesion CA-Containing Neurons in Neonatal Rats

In 1972, Breese and Traylor administered 6-OHDA into the cisterna magna of developing rats to determine whether the 6-OHDA would destroy CA-containing neurons during this period (Breese and Traylor 1972). They found that this procedure produces a marked reduction in brain DA, NE, and tyrosine hydroxylase activity (TH; the rate limiting enzyme required for DA synthesis) (Breese and Traylor 1972). In subsequent studies to reduce only brain



**Fig. 1** Tyrosine hydroxylase immunohistochemistry after neonatal 6-OHDA lesioning in rats. Immunohistochemical detection of tyrosine hydroxylase in coronal sections representing the striatum, nucleus accumbens, and medial prefrontal cortex of adult rats after neonatal sham (*top*) and 6-OHDA (*bottom*) lesioning

DA, selective destruction of DA neurons was achieved by pretreating rats with desipramine prior to injection of 6-OHDA either into the cisterna magna or into the cerebral ventricles (Smith et al. 1973). At 100–150 µg doses of 6-OHDA, injection into either of these ventricular sites was found to produce a major destruction of DA- and NE-containing neurons (>95 %) within 72 h (Breese et al. 1984a, 1985a, b, 1987a; Berger et al. 1985; Snyder et al. 1986). In fact, as early as 4 days postinjection into the ventricles, DAergic cell bodies and dendrites were reported to disappear entirely from the SN (Fernandes Xavier et al. 1994) and cause a considerable reduction in number in the VTA (Fernandes Xavier et al. 1994; Breese et al. 1984a, 1985a, b, 1987a; Berger et al. 1985; Snyder et al. 1986). Consequently, the DAergic terminal fibers in the striatum are drastically reduced by 6-OHDA (Snyder et al. 1986; Descarries et al. 1992), as are neurons in certain target regions of the VTA, including the septum, nucleus accumbens, and frontal cortex (Luthman et al. 1990a). This extensive DAergic denervation explains the profound loss of brain DA content measured in these animals during adulthood (Breese and Traylor 1972; Breese et al. 1984a; Stachowiak et al. 1984; Bruno et al. 1987; Jackson et al. 1988; Dewar et al. 1990). In Fig. 1, the extent of DA depletion in the striatum and frontal cortex of adult rats lesioned with 6-OHDA as neonates is revealed through the immunohistochemical detection of TH. This perinatal loss of DA in neonatal rats has been the focus of investigations aimed at understanding how this loss relates to the pathophysiology of the developmental brain disorder LNS in which DA is reduced during development (Breese et al. 2005).

### 3 Behavioral Responses After 6-OHDA Lesions

### 3.1 Behavioral Impairment and Recovery of Function in Rats Lesioned as Adults

As noted earlier, extensive bilateral lesions of nigrostriatal DA neurons in adult rats produce a syndrome that is analogous to PD in humans (Marshall et al. 1974; Ungerstedt 1971a; Zigmond and Stricker 1984). The rats exhibit symptoms such as bradykinesia/akinesia, rigidity, sensorimotor neglect, excessive bracing and clinging reactions, festination, aphagia, adipsia, and cognitive dysfunction (Marshall 1979; Stricker and Zigmond 1984; Ungerstedt 1971a; Zigmond and Stricker 1972; Zis et al. 1974). The occurrence and severity of these behavioral deficits appears to be strongly correlated with the degree of DA depletion in the striatum (Lees et al. 1985). For example, adult rats with essentially total bilateral depletions of striatal DA (>95 %) can perish as a consequence of these behavioral impairments if they are not tube-fed intragastrically with nutrients. In spite of this tube-feeding effort over an extended period, the rats rarely regain normal ingestive behavior and continue to show sensorimotor neglect and profound motor and cognitive impairments (Marshall 1979: Zigmond and Stricker 1973). These 6-OHDA-lesioned adult rats continue to suffer from residual regulatory deficits, particularly in their ability to respond to stress and acute homeostatic imbalances (Snyder et al. 1985; Stricker and Zigmond 1974; Marshall and Teitelbaum 1974). In fact, when challenged with an exposure to stressor, the severe ingestive and motor impairments can be reinstated in these animals to the level of severity seen acutely after the lesion (Robinson et al. 1990). Adult rats with extensive, yet less severe, bilateral 6-OHDA lesions of striatal DA (80–90 %) initially display behavioral deficits (Baez et al. 1977; Fibiger et al. 1973; Spirduso et al. 1985) but show significant recovery over time (Altar et al. 1987; Creese et al. 1977; Dravid et al. 1984; Neve et al. 1982; Nisenbaum et al. 1986). Similarly, patients with PD manifest clinical symptoms only after a loss of 80 % of the DAergic input into the striatum (Bernheimer et al. 1973; Fearnley and Lees 1991). Therefore, while nigrostriatal DAergic activity is required for normal behavioral function, only 10-20 % of nigrostriatal DA input is actually needed to maintain fairly normal function in both rats and humans.

### 3.2 Sparing of Behavioral Impairment and Recovery of Function in Rats Lesioned as Neonates

The behavior of rats lesioned with 6-OHDA as neonates is very different from that of rats lesioned with 6-OHDA during adulthood. Rats lesioned with 6-OHDA as

neonates do not display any outward symptoms of aphagia or adipsia associated with adult bilateral lesions (Breese and Traylor 1972; Smith et al. 1973). In fact, while these neonatal lesioned rats are smaller in stature (Breese and Traylor 1972), they show few overt behavioral deficits that distinguish them from non-lesioned rats during development or in subsequent adulthood, even when their DA levels are depleted by 99 % (Breese et al. 1984a; Bruno et al. 1984; Erinoff et al. 1979; Smith et al. 1973). Near-total depletion of striatal DA incurred prior to postnatal day 27 (P27) produces no obvious deficits in sensorimotor function either soon after the lesion or after the rats reach adulthood (Weihmuller and Bruno 1989a), nor is there any evidence of profoundly impaired ingestive or motor function (Potter and Bruno 1989; Weihmuller and Bruno 1989b). In early studies, Breese and Traylor showed that neonatal rats with reductions in both DA and NE remained virtually indistinguishable from non-lesioned controls except for a slight decrease in body weight and a lack of self-grooming (Breese and Traylor 1972). Smith and colleagues later showed that the reduced body weight induced by the 6-OHDA reduction in food consumption was related to reduced DA and not to a loss of NE (Smith et al. 1973). Although neonatal rats with DA or with both CA-containing neurons destroyed exhibit reduced body weight (Breese and Traylor 1972; Smith et al. 1973), they are still able to suckle and grow and can be weaned readily onto solid food and water as early as 27 days of age (Bruno et al. 1984, 1986; Weihmuller and Bruno 1989b; Smith et al. 1973).

The apparent sparing of behavioral function in neonatal lesioned rats, however, is not a complete one. Deficiencies in rats with DA selectively reduced during development were eventually uncovered with rigorous behavioral assessments of challenges to functional measures (Smith et al. 1973). For example, Smith and colleagues found that neonatal lesioned rats with selective reduction of brain DA do not ingest sucrose even though intake of water was normal (Smith et al. 1973), whereas rats with only brain NE reduced exhibit an increase in the sucrose drinking over that of controls. Others found that the neonatal lesioned rats with DA reduced do not eat in response to acute glucoprivation and do not drink water or saline in response to hypovolemia (Bruno et al. 1986). These deficits in response to acute homeostatic imbalances are similar to those seen in rats with adult lesions (Bruno et al. 1986). Further, studies showed that neonatal lesioned rats with only DA reduced exhibit a total deficiency in acquiring the shuttle-box avoidance response in adulthood, while rats with only NE reduced exhibit an elevated acquisition in this task (Smith et al. 1973). As adults, the neonatal lesioned rats show significant deficits in classical and aversive conditioning (Raskin et al. 1983; Whishaw et al. 1987), escape and avoidance learning (Shaywitz et al. 1976b; Pappas et al. 1980), spatial navigation (Whishaw et al. 1987; Pappas et al. 1992), and independent limb/ tongue use and limb posture (Whishaw et al. 1987), suggestive that the neonatal DA depletion produces lasting impairments in learning ability and skilled motor control. Rats depleted of DA as neonates also show deficits in motoric capacity for high-rate operant responding (lever-pressing) during intracranial self-stimulation to the lateral hypothalamus (Stellar et al. 1988) and VTA (Takeichi et al. 1986), as well as acquisition deficits in operant responding for water and other reinforcing stimuli (Heffner and Seiden 1983; Moy 1995), just as do adult lesioned rats (Cooper and Breese 1975). Adult rats with DA destroyed as neonates with 6-OHDA also exhibit reduced locomotion, whereas those that had NE selectively reduced exhibit a nearly doubling of locomotion (Smith et al. 1973). Other reports have shown that animals lesioned with 6-OHDA during development exhibit hyperactivity in adolescence similar to patients with ADHD (Shaywitz et al. 1976b; Miller et al. 1981; Luthman et al. 1989a). Thus, while rats with neonatal DA-specific lesions cope quite well in the absence of challenges, exposure of these animals to such confrontations clearly reveals that severe functional deficiencies persist.

### 4 Compensatory Neural Responses After 6-OHDA Lesions

### 4.1 Adaptations in the DA System of Adult Lesioned Rats

In both neonatal and adult 6-OHDA-lesioned rats, a number of compensatory responses have been identified in the striatal DA system that could be involved in counterbalancing the effects of the lesion. While adaptive changes related to 6-OHDA lesioning likely occur when the toxin destroys NE-containing neurons, this following section focuses only on adaptations that occur after the selective lesioning of DA neurons with 6-OHDA. The adaptive changes in the DA system that occur in response to 6-OHDA lesioning are largely neurochemical, involving presynaptic changes in the remaining DA terminals as well as postsynaptic changes in target cells that are dependent upon lesion size. Such changes at synapses likely enable the spared DA neurons to upregulate their activity in order to regain functional control of their targets (Zigmond 1997). Initially, the adaptations observed in adult lesioning of DA neurons.

Presynaptic DA Changes. In adult rats receiving 6-OHDA lesions of the nigrostriatal DA pathway, presynaptic changes occur that are largely related to an enhanced metabolism of surviving DA neurons. Well established is that the remaining DA neurons greatly increase their rate of DA synthesis at striatal terminals (Agid et al. 1973; Hefti et al. 1980, 1985; Wolf et al. 1989), a finding attributed to the ability of these cells to upregulate their intrinsic TH activity presumably to meet the increased demand for releasable DA (Zigmond et al. 1984). In fact, a direct correlation exists between the degree of DA depletion by 6-OHDA and the extent of upregulation of TH activity in the spared terminals, an increase which in most cases can be greater than sixfold (Zigmond et al. 1984). Using intracerebral microdialysis techniques in adult rats with unilateral DA lesions, Robinson and Whishaw showed that the extracellular concentrations of DA in the denervated striatal hemisphere are much higher than predicted given the extent of DA depletion measured in the postmortem tissue (Robinson and Whishaw 1988). Later microdialysis studies in adult rats with bilateral 6-OHDA lesions showed that the extracellular DA concentrations remain close to normal even when the DA destruction in both striatal hemispheres was nearly complete (Castaneda et al. 1990b). Additionally, an increase in the rate of firing and burst firing activity of the remaining striatal DA neurons occurs (Hollerman and Grace 1990), which together can contribute to the increased levels of DA release as well as to the amount of DA release per action potential in the lesioned striatum (Stachowiak et al. 1987). At the same time, a decrease in DA reuptake, produced by the destruction of DAT on DA terminals by 6-OHDA, contributes to a slower rate of DA clearance from the extracellular space which, in combination with the disappearance of presynaptic DA autoreceptors, results in an increased diffusibility of DA throughout the lesioned striatum (Zigmond et al. 1984; Snyder et al. 1990).

An additional contributing factor to the functional sparing or recovery related to presynaptic function seen after extensive 6-OHDA lesions in adult rats appears to relate to sprouting of remaining DA neuronal terminals into denervated striatal regions. This growth occurs from an undamaged axon, possibly a neighboring axon in the same tract or an axon from a different tract, to fill a vacated adjacent space. Although the nigrostriatal DA projection is predominately ipsilateral, there is also a minor crossed projection that arises from the contralateral SN (Fallon and Loughlin 1982; Gerfen et al. 1982; Morgan et al. 1986). While DA fibers of the ipsilateral projection have a clear capacity to undergo this regenerative sprouting (Bohn et al. 1987; Hansen et al. 1995; Onn et al. 1986), Pritzel and colleagues used a retrograde labeling technique to show an increased number of contralaterally labeled nigral neurons that innervated the striatum 7 days after an extensive unilateral lesion of the ipsilateral nigrostriatal bundle in adult rats (Pritzel et al. 1983). This sprouting of DA neurons to reinnervate the DA-denervated striatum could be a possible basis for the eventual recovery of unilaterally lesioned adult rats from their initial turning posture after exposure to the 6-OHDA. Ultrastructural tracer studies have also revealed the appearance of hypertrophic DAergic fibers and growth-cone-like structures and the expansion of a terminal arbor network over a period of 1-4 months after the adult unilateral 6-OHDA lesioning (Blanchard et al. 1996; Liberatore et al. 1999; Finkelstein et al. 2000). Evidence suggests that this process is regulated by both supersensitive postsynaptic receptors and glial cells activated by the lesion (Batchelor et al. 1999; Parish et al. 2002; Tripanichkul et al. 2003). It has been hypothesized that the enlargement of synaptic varicosities and terminal arborizations observed after adult lesions could help compensate for the loss of DA by delivering larger boluses of DA to the denervated striatum, especially after presynaptic DA reuptake mechanisms and autoreceptors have been disrupted by the 6-OHDA lesion (Castaneda et al. 1990b; van Horne et al. 1992). Studies with DA-rich fetal SN or adrenal medulla transplants showed that replacement of only a very small fraction of the normal complement of DA is sufficient to restore some functional control of the striatum (Carder et al. 1987, 1989; Snyder-Keller et al. 1989).

Collectively, these studies suggested that the presynaptic adaptations in the remaining DA terminals likely exist to normalize the extracellular DA concentrations in the denervated striatum – changes which contribute to the recovery of function in the 6-OHDA-lesioned rats (Robinson and Whishaw 1988; Robinson et al. 1990, 1994; Abercrombie et al. 1990; Castaneda et al. 1990b; Parsons et al. 1991; Touchet and Bennett 1989). However, when striatal DA depletion exceeds 95 %, a marked

drop in extracellular levels of DA occurs, presumably reducing drastically the degree of behavioral function because of the extensive loss of presynaptic DA adaptation (Abercrombie et al. 1990; Castaneda et al. 1990b; Robinson et al. 1990). Therefore, at least 5 % of the DAergic cells must be present to produce an adequate supply of extracellular DA to maintain control over striatal targets and ultimately to sustain some function in the adult lesioned animals (Robinson et al. 1990).

*Postsynaptic DA Changes*. In adult 6-OHDA-lesioned rats, postsynaptic changes are manifested when the degree of striatal DA depletion by 6-OHDA becomes more extreme – an additional adaptation that is likely to contribute to the recovery process induced by the lesioning (Zigmond 1997; Zigmond and Stricker 1980; Staunton et al. 1981). This recovery process can enable the lesioned animals to amplify a weak DA signal (Zigmond 1997). A number of quantitative autoradiography and small animal positron tomography (PET) studies have consistently reported an increase in the density of postsynaptic D<sub>2</sub>-DA receptors in the striatum of rats with extensive 6-OHDA lesioning of DA during adulthood (Creese et al. 1977; Savasta et al. 1987; Staunton et al. 1981; Murrin et al. 1979; Neve et al. 1982; Graham et al. 1990; Przedborski et al. 1995; MacKenzie et al. 1989; Joyce 1991a; Nikolaus et al. 2003; Hume et al. 1995; Nguyen et al. 2000). While the majority of these studies have been conducted in rats with unilateral DA depletions, a similar increase has been found in rats with bilateral loss of 90 % or more of total DA content in both striatal hemispheres (MacKenzie et al. 1989; Joyce 1991a). Autoradiography studies using  $[{}^{3}H]$  spiroperidol to localize the changes in D<sub>2</sub> receptors revealed increases in the lateral parts of the striatum after short survival times (Savasta et al. 1987), with a more diffuse upregulation into medial parts by 7 months after such lesions (Chritin et al. 1992). Other studies showed that the increase in  $D_2$ receptor density is accompanied by an upregulation in D<sub>2</sub> receptor mRNA levels (Gerfen et al. 1990; Angulo et al. 1991; Neve et al. 1991; Cadet et al. 1992; Lisovoski et al. 1992; Chritin et al. 1993). Brene and colleagues found a  $D_2$  receptor mRNA increase on medium-sized  $\gamma$ -aminobutyric acid (GABA)-containing neurons in the lateral striatum 3 weeks after a unilateral 6-OHDA lesion (Brene et al. 1990). Reports have suggested that the increase in  $D_2$  receptors is due to a propensity of mRNA for the D<sub>2</sub>-long spliced variant (Martres et al. 1992). Interestingly, this isoform of the D<sub>2</sub> receptor is preferentially involved in postsynaptic responses to DA under normal conditions (Usiello et al. 2000; Wang et al. 2000). Consistent with the changes in  $D_2$  receptor synthesis, the recovery rate of  $D_2$ receptor binding following irreversible blockade with N-ethoxycarbonyl-2-ethoxy-1,2-dihydroquinoline (EEDQ) was found to be enhanced in lesioned striatum 4-5 weeks after unilateral 6-OHDA treatment in adult rats (Neve et al. 1985). This increase in recovery was attributed to an increase in the rate of reappearance of  $D_2$  receptors into synaptic membranes. Irrespective of these findings, the physiological consequences of  $D_2$  receptor upregulation on striatal function remain unknown, as increased  $D_2$  receptor density was found to have no effect on the inhibitory responsiveness of spontaneously firing striatal neurons after exposure to iontophoresed DA or to the D2 receptor agonist 2-(N-phenyl-ethyl-N-propyl)amino-5-hydroxytetralin in adult lesioned animals (Radja et al. 1993b). Nonetheless, this

adaptive increase in  $D_2$  receptors could provide another means by which to amplify a weak DA signal induced by the DA lesioning (Zigmond 1997).

In adult lesioned rats, the effects of 6-OHDA-induced destruction of DA neurons on the regulation of other DA receptor subtypes, in particular the  $D_1$  receptor, remain a contentious issue (Reader and Dewar 1999). Studies have reported that the density of  $D_1$  receptors is either increased (Buonamici et al. 1986; Fornaretto et al. 1993; Narang and Wamsley 1995), unaltered (Altar and Marien 1987; Savasta et al. 1988; Lawler et al. 1995), or decreased in the striatum of these animals (Marshall et al. 1989; Stromberg et al. 1995). Further studies have reported either no change (Brene et al. 1990; Fornaretto et al. 1993; Narang and Wamsley 1995) or a slight decrease in D<sub>1</sub> mRNA levels after such lesions (Stromberg et al. 1995). Such variability might be due to (1) the amount of damage induced by the adult 6-OHDA lesion (Iwata et al. 1996; Reader and Dewar 1999), (2) the period of survival following the lesion (Fornaretto et al. 1993; Narang and Wamsley 1995), and/or (3) the site and conditions selected for the 6-OHDA treatment (Narang and Wamsley 1995; Reader and Dewar 1999). Therefore, parameters such as time course and spatial distribution of the lesion must be considered when comparing and interpreting the various changes reported for D<sub>1</sub> receptor function after 6-OHDA lesions in adult rats.

### 4.2 Adaptations in the DA System of Neonatal Lesioned Rats

Presynaptic DA Changes. While adult 6-OHDA-lesioned rats remain exclusively dependent upon the amount of DA that remains after 6-OHDA treatment, rats lesioned as neonates survive quite well and continue to exhibit sparing of function even when only 1-2 % of the nigrostriatal DA neurons remain. In this respect, an adaptive change which appears to be unique to neonatal 6-OHDA-lesioned rats includes an increased ability of post-lesion residual neurons to release DA, even when 99 % of the DA content in both striatal hemispheres is lost (Carder et al. 1989; Castaneda et al. 1990b). The few surviving DA neurons after neonatal lesions undergo presynaptic metabolic changes that persist well into adulthood and are similar to those seen after adult lesions (Molina-Holgado et al. 1994). These changes include an increase in the synthesis of DA, suggestive of an enhanced release of DA from spared DA terminals (Molina-Holgado et al. 1994). As with adult lesions, it has been suggested that these presynaptic adaptations in the neonatal lesioned rats account for the maintenance of a relatively high extracellular DA concentration in the denervated striatum during the resting state, as measured by intracerebral microdialysis (Castaneda et al. 1990a). These higher levels of extraneuronal DA are quite surprising given the meager 1-2 % survival rate of striatal terminals after such lesions. It has been hypothesized that the storage capacity of the remaining terminals may be increased to protect DA from intraneuronal oxidative deamination (Reader and Dewar 1999), particularly since studies have shown that the actual size of the terminals becomes larger (Pickel et al. 1992). An alternative hypothesis is that the membrane properties of the DA-containing vesicles may become altered to facilitate the increased transmitter release (Reader and Dewar 1999). Regardless, the normalization of extracellular DA levels in the denervated striatum is likely a crucial component in the sparing of behavioral function seen in rats after destruction of neonatal DA neurons.

When neonatal lesioned rats are administered an additional 6-OHDA treatment in adulthood to further reduce presynaptic function, these animals show symptoms of aphagia, adipsia, and akinesia similar to adult lesioned rats, suggesting that the residual DA activity is involved in the sparing of these particular functions subsequent to the neonatal lesion (Rogers and Dunnett 1989b). Likewise, neonatal lesioned rats are hypersensitive to the disruption of residual DA function with the catecholamine synthesis inhibitor  $\alpha$ -methyl-para-tyrosine ( $\alpha$ -MT) that results in deficits in feeding as well as other behaviors (Rogers and Dunnett 1989a), just as observed in adult 6-OHDA-lesioned rats (Cooper et al. 1972, 1973b). Collectively, such outcomes provide convincing evidence that the residual DA neurons are necessary for functional sparing in these neonate lesioned animals.

*Postsynaptic DA Receptor Changes*. In rats lesioned as neonates, early studies reported no changes in the binding of [<sup>3</sup>H]-spiperone to D<sub>2</sub> receptors after i.c. administration of 6-OHDA (Breese et al. 1987a; Duncan et al. 1987; Kostrzewa and Brus 1991; Kostrzewa and Hamdi 1991). However, subsequent studies in lesioned rats using [<sup>3</sup>H]-raclopride demonstrated significant increases of 30–40 % in D<sub>2</sub> receptor density over that in controls (Dewar et al. 1990). This change occurred in the rostral portion of the striatum between 1 and 3 months after the 6-OHDA lesion (Dewar et al. 1990), but increased throughout the entirety of the striatum at a later time after the lesion (Radja et al. 1993b). Because studies showed that the recovery of D<sub>2</sub> binding after in vivo receptor alkylation with EEDQ was unaltered, this D<sub>2</sub> receptor proliferation after the neonatal lesioning was proposed to be the result of posttranscriptional modifications (Dewar et al. 1997).

Unlike adult 6-OHDA-lesioned animals, the increase in density of the  $D_2$ receptors in the bilaterally lesioned neonates is not accompanied by a sustained increase in mRNA levels (Chen and Weiss 1991; Radja et al. 1993b). In contrast to this finding, unilateral lesions of the nigrostriatal DA pathway in neonatal rats were found to produce a 20-25 % increase in the density of D<sub>2</sub> receptors (radiolabeled with the D<sub>2</sub> antagonist quinpirole, LY 171555) which was accompanied by an upregulation of  $D_2$  mRNA levels in the rostral striatum 9 months after the lesion (Mennicken et al. 1995). This latter finding is consistent with the increase in the density of  $D_2$  receptors being related to an increase in the synthesis of the  $D_2$ receptor. Further, since binding in a control was the same as the contralateral side to the lesion, receptor changes were not likely related to adaptation on the unlesioned side of the brain (Mennicken et al. 1995). The increase in  $D_2$  receptor binding and mRNA levels was not observed in similarly lesioned rats that received DA-rich grafts intrastriatally 5 days after the 6-OHDA lesion (Mennicken et al. 1995), a finding supportive of the view that the adaptive increase in  $D_2$  receptors was related to the destruction of the striatal DA neurons.

Similar to the  $D_1$  receptor binding studies in adult 6-OHDA-lesioned rats, there are conflicting reports of slight increases (Broaddus and Bennett 1990), slight

decreases (Dewar et al. 1990; Dewar et al. 1997; Molina-Holgado et al. 1995), large decreases (Gelbard et al. 1990), or no change at all (Breese et al. 1987a; Luthman et al. 1990b) in the density of striatal  $D_1$  receptors in neonatal 6-OHDA-lesioned rats. Also, there is no change in the percentage of low-affinity ( $R_1$ ) and high-affinity  $(R_{\rm H})$  binding sites for striatal D<sub>1</sub> receptors in adult rats that were lesioned as neonates with 6-OHDA (Gong, Kostrzewa et al. 1994). So far, studies attempting to explore changes in mRNA encoding for the D<sub>1</sub> receptor in these neonatal lesioned rats have failed to find any significant changes in these rats compared to controls (Duncan et al. 1993). In contrast to the latter findings, Gong et al. found that D<sub>1</sub> receptor mRNA is reduced by 25–50 % in neonatally 6-OHDA-lesioned rats at 6 weeks post-birth and that 4 weekly treatments with the  $D_1$  receptor agonist SKF 38393 restore  $D_1$  receptor mRNA to control levels (Gong et al. 1994). It is also important to note here that there are at least six different forms of the DA receptor, and the terms  $D_1$  and  $D_2$  refer more to subfamilies of receptor type and not to a single DA receptor isoform (Strange 1993). The  $D_1$  class of DA receptors, which consists of the  $D_1$  and  $D_5$  subtypes, stimulates adenylate cyclase (AC) activity, while the  $D_2$  class, comprised of the  $D_2$  short,  $D_2$  long,  $D_3$ , and  $D_4$  subtypes, generally inhibits it. It is possible that some of the compensation seen after 6-OHDA lesioning in rats is related to changes in DA receptor types other than  $D_1$  and  $D_2$  receptors, although such changes could also facilitate some of the behavioral disturbances seen in neonatal lesioned animals. Specifically, Zhang and colleagues found an increase in the density of  $D_4$  receptors in the striatum and a decrease in the nucleus accumbens at P25, but not at P37 or P60 after neonatal 6-OHDA lesions (Zhang et al. 2002a, b), with only minor changes in  $D_1$  and  $D_2$ receptor binding in various forebrain regions. The change in D<sub>4</sub> receptors, but not in  $D_1$  and  $D_2$  receptors, was positively correlated with the time course of motor hyperactivity exhibited by the neonatal lesioned animals, suggesting that  $D_4$  receptors may play a pivotal role in this response (Zhang et al. 2002a, b). In subsequent studies, Zhang and colleagues also showed that the administration of D<sub>4</sub> antagonists L-745,870 and U-101,958 dose-dependently inhibits the motor hyperactivity in neonatal lesioned animals (Zhang et al. 2002a).

### 4.3 Adaptations in the 5-HT System by 6-OHDA Lesions

*5-HT Hyperinnervation*. An alternative view to the direct role of residual DA and receptor changes in the sparing or recovery of function in rats after 6-OHDA lesions is that compensatory mechanisms within the 5-HT system might provide some replacement for the loss of DA function. In this respect, following lesioning of DA neurons in neonatal animals, an extensive hyperinnervation of 5-HT of axon terminals or varicosities arising from the dorsal raphe nucleus is found in the striatum that is accompanied by an increase in 5-HT levels (Breese et al. 1984a; Berger et al. 1985; Luthman et al. 1987; Snyder et al. 1986; Stachowiak et al. 1984; Towle et al. 1989; Descarries et al. 1992; Molina-Holgado et al. 1994). So long as 80 % of the striatal DA is lost, both unilateral and bilateral DA depletions can

produce this 5-HT hyperinnervation in neonatal lesioned rats (Towle et al. 1989; Gong et al. 1993b; Penit-Soria et al. 1997). In rats lesioned at 3 days of age, this extent of DA loss can produce a severalfold increase in 5-HT content in the rostral striatum 2–5 months post-lesioning, along with an upregulation in high-affinity 5-HT uptake (Breese et al. 1984a; Stachowiak et al. 1984) and 5-HT transporter density (Molina-Holgado et al. 1994). While initially reported that this adaptive 5-HT hyperinnervation was not seen in adult rats after a comparable bilateral DA lesion of the striatum (Breese et al. 1984a), striatal 5-HT hyperinnervation was later reported to occur after unilateral 6-OHDA lesions in adult rats if striatal DA content was severely reduced by over 95 % (Zhou et al. 1991; Guerra et al. 1997; Maeda et al. 2003).

Apparently, the trigger for this proliferation of 5-HT-containing fibers is the extensive loss of striatal DA neurons (Kostrzewa et al. 1993; Towle et al. 1989) even though the loss of DA occurs several days before the development of 5-HT innervation (Lidov and Molliver 1982). Further evidence in accord with this view is that the 5-HT terminals do not occupy synaptic sites vacated by the loss of DA neurons and do not interfere with DA outgrowth arising from transplanted DA-rich cells derived from the ventral mesencephalon (Snyder-Keller et al. 1989; Abrous et al. 1993a). Additionally, replacement of lost DA innervation by transplanted DArich cells shortly after the lesion prevents the 5-HT hyperinnervation (Snyder-Keller et al. 1989; Abrous et al. 1993a). The sprouted axons that accumulated significant amounts of 5-HT (Molina-Holgado et al. 1994) tripled their extracellular 5-HT levels under stimulatory release conditions (Jackson and Abercrombie 1992). However, intracerebral microdialysis studies under resting conditions demonstrated that the striatal 5-HT hyperinnervation is not associated with enhanced levels of basal extracellular 5-HT, probably due to an increase in high-affinity 5-HT uptake (Jackson and Abercrombie 1992). Further, Descarries et al. revealed that the hyperinnervation is not accompanied by an increase in the number of synaptic contacts made by the 5-HT varicosities (Descarries et al. 1992). On the other hand, Jackson and colleagues found that that endogenously released 5-HT could affect acetylcholine (Ach) release in neonatally denervated striatum (Jackson et al. 1988). Additionally, 5-HT release could exert effects on striatal DA terminals through indirect diffusion processes (Descarries and Mechawar 2000; Umbriaco et al. 1995). For example, the release of 5-HT from hyperinnervated fibers could contribute to enhancing DA release from residual DA terminals by entry into the presynaptic DA terminals, which could in turn facilitate DA release from these remaining neurons (Navailles et al. 2010).

While initially hypothesized that the increased striatal 5-HT might be responsible for the sparing phenomenon in neonatal DA-lesioned animals (Towle et al. 1989), а combination of 6-OHDA with the 5-HT neurotoxin 5.7dihydroxytryptamine (5,7-DHT) (Breese et al. 1978) given to 3-day-old rat pups to lesion both 5-HT and DA neurons produced no greater behavioral dysfunction than 6-OHDA lesions alone (Bruno et al. 1987). Consequently, if indeed 5-HT was serving as a substitute for the lost DA, then it would be expected that rats with co-lesions of the two transmitter systems would exhibit severe impairments,

perhaps similar to rats lesioned in adulthood. Further, behavioral sparing was still observed in rats treated with 6-OHDA at P15 and P27 (Bruno et al. 1987), even though the neurochemical effects of 6-OHDA lesions given at this point in development do not include the elevated 5-HT levels typical of early lesioning of neonatal DA neurons (Bruno et al. 1987). Given these various findings, the 5-HT hyperinnervation does not appear to be a critical component in the functional sparing seen in neonatal 6-OHDA-lesioned rats.

5-HT Receptor Binding. In adult rats, the effect of 6-OHDA lesions on various 5-HT receptors and their mRNA levels in the brain is controversial, as a number of receptor autoradiographic and in situ hybridization studies have provided conflicting results. Early studies suggested that 6-OHDA-induced DA depletion in adult rats does not alter either 5-HT receptor densities or the inhibitory responsiveness of striatal neurons to iontophoretic application of 5-HT or its agonists (el Mansari et al. 1994). Despite this reported lack of effect of DA destruction on 5-HT receptor function, later studies found an increase in 5-HT<sub>2A</sub> receptor mRNA levels in the striatum after unilateral 6-OHDA lesions in adult rats (Numan et al. 1995; Zhang et al. 2007) in which 5-HT is elevated (Zhou et al. 1991; Guerra et al. 1997; Maeda et al. 2003). This striatal 5-HT receptor increase was abolished by L-dopa treatment (Zhang et al. 2007), suggestive that the regulation of striatal 5-HT<sub>2A</sub> receptors is primarily dependent upon DA acting on DA receptors. At odds with these results, a more recent study performed in adult 6-OHDA-lesioned animals using [<sup>3</sup>H] ketanserin-binding autoradiography showed reduced 5-HT<sub>2A</sub> receptor levels in the striatum, as well as in the cingulate, insular, prefrontal, and primary somatosensory cortices (Li et al. 2010). In contrast to this increase, studies in adult 6-OHDA-lesioned rats have also shown a decrease in 5-HT<sub>2C</sub> receptor mRNA levels in the striatum (Zhang et al. 2007). L-dopa treatment had no effect on 5-HT<sub>2C</sub> mRNA levels in this region (Zhang et al. 2007), suggesting that the 5-HT<sub>2C</sub> and 5-HT<sub>2A</sub> receptors are differentially regulated by the nigrostriatal DA function. Also in adult 6-OHDA-lesioned animals, one study found no change in striatal 5-HT<sub>4</sub> receptor levels upon binding with the selective radioligand [<sup>125</sup>I] SB 207710 (Patel et al. 1995), while another found increases in the caudal, but not rostral, striatum and globus pallidus upon binding with the 5-HT<sub>4</sub> receptor antagonist [<sup>3</sup>H] GR113803 (Compan et al. 1996). Such discrepancies could be due to methodological factors, such as the type of ligand used, the state of receptor affinity measured, the actual placement of the 6-OHDA lesion, and/or the part of the brain region analyzed.

In striatum of neonatal 6-OHDA-lesioned rats in which 5-HT was elevated, quantitative ligand binding utilizing autoradiography has revealed changes in several 5-HT receptor subtypes at adulthood (Radja et al. 1993a; Reader and Dewar 1999; Kostrzewa et al. 1998; Laprade et al. 1996). Binding of [<sup>3</sup>H] 5-HT revealed an increase throughout the striatum and SN after neonatal lesioning, but not in the globus pallidus (Radja et al. 1993a), suggestive that this upregulation occurs primarily on striatonigral as opposed to striatopallidal projection neurons. Labeling of 5-HT<sub>1B</sub> receptors with the nonselective radioligand [<sup>125</sup>I] cyanopindolol revealed an increase in the binding density of these receptors

throughout the striatum (i.e., both rostral and caudal halves), globus pallidus, and SN 3 months after the neonatal lesions (Radja et al. 1993a). Since previous studies showed that 5-HT<sub>1B</sub> mRNA was present in the striatum but not in SN or globus pallidus of rats (Bruinvels et al. 1994), the increase in 5-HT<sub>1B</sub> binding seen in the striatum, globus pallidus, and SN was suggested to result from an upregulation and increased transport of these receptors on striatal GABAergic projection neurons (Reader and Dewar 1999). Moreover, labeling of 5-HT<sub>2A</sub> receptors with [<sup>125</sup>I] iodimethoxyphenylamino-propane, or [<sup>125</sup>I] DOI, revealed a marked increase in this receptor subtype in the rostral striatum of adult rats lesioned as neonates (Radja et al. 1993a). The anatomical distribution of increased 5-HT<sub>2A</sub> receptors strongly correlated with that of 5-HT hyperinnervation in the rostral striatum, initially suggestive of a relationship between the two phenomena. However, studies established that the majority of 5-HT<sub>2A</sub> receptors are postsynaptic to 5-HT neurons (Mengod et al. 1990) and that the increase in 5-HT<sub>2A</sub> receptors occurs independent of the 5-HT hyperinnervation (Laprade et al. 1996; Basura and Walker 1999; Fox and Brotchie 2000; Kostrzewa et al. 1998). The increased 5-HT<sub>2A</sub> binding density in the rostral striatum was also associated with a considerable increase in mRNA encoding for this receptor within lateral parts of the striatum of the neonatal lesioned rats (Laprade et al. 1996). In these neonatal lesioned animals, both changes were abolished after chronic treatment with apomorphine or the  $D_1$  receptor agonist SKF-38393, suggesting that  $D_1$  receptor activation participates in the degree of 5-HT<sub>2A</sub> receptor expression (Laprade et al. 1996). Examination of the 5-HT<sub>1A</sub> receptor subtype upon specific labeling with [<sup>3</sup>H] 8-hydroxy-dipropylaminotetralin or [<sup>3</sup>H] 8-OH-DPAT revealed no changes in the binding density of these receptors in adult striatum after neonatal 6-OHDA lesions (Radja et al. 1993a), nor was there any changes in 5-HT<sub>1A</sub> mRNA levels (Numan et al. 1995).

# 4.4 Adaptations in GABA and Other Amino Acid Transmitter Systems by 6-OHDA

Discrete populations of GABAergic medium-sized spiny neurons in the striatum project their axons to the globus pallidus, entopeduncular nucleus (internal pallidum), and SN (Beckstead and Cruz 1986; Bolam and Smith 1990; Loopuijt and van der Kooy 1985). The inhibitory activity of these GABAergic neurons (van der Heyden et al. 1980) appears to be modulated by the ascending nigrostriatal DA fibers arising from the SNpc (van der Heyden et al. 1980; Samuel et al. 1988; Vernier et al. 1988; Reid et al. 1990). This efferent GABAergic population from the striatum is predominately subdivided into two subclasses, defined by the nature of their co-transmitters (Penny et al. 1986; Reiner and Anderson 1990) and the differential expression of  $D_1$  and  $D_2$  receptor subtypes on their surface. The striatal GABAergic neurons that project to the SN and entopeduncular nucleus mainly express  $D_1$  receptors and their axons contain the neuropeptides substance P, neurokinin A, and dynorphin, while those that project to the globus pallidus predominately express  $D_2$  receptors and contain the neuropeptide enkephalin

(Gerfen and Young 1988; Gerfen et al. 1990, 1991; Gerfen 1992). These striatonigral and striatopallidal pathways work in concert to regulate thalamic output to motor areas of the cortex and thus play an integral role in the execution of movement. More recently, a third, smaller population of striatal GABAergic neurons has been identified that co-expresses both  $D_1$  and  $D_2$  receptors as well as the neuropeptides substance P and enkephalin (Perreault et al. 2011). Unlike the striatal GABAergic neurons expressing only  $D_1$  or  $D_2$  receptors, these  $D_1/D_2$ receptor striatal projection neurons terminate in regions within both the striatonigral and striatopallidal pathways, as well as regions containing DAergic cell bodies such as the VTA (Deng et al. 2006; Wang et al. 2006, 2007). While little is known about the functional relevance of these latter neurons, they are thought to contribute to the regulation of thalamic neurotransmission, perhaps with the purpose of maintaining homeostatic balance between the striatonigral and striatopallidal pathways (Perreault et al. 2011). An imbalance between the striatonigral and striatopallidal pathways, such as seen with DAergic degeneration in humans and in animals (Albin et al. 1989; DeLong 1990; Hikosaka et al. 2000; Bezard et al. 2001; Obeso et al. 2004), can affect the final outflow to the thalamus and contribute to motor disturbances (Gerfen et al. 1990; Gerfen 2000; Mallet et al. 2006).

Increased GABA Levels. In adult rats, unilateral DA depletion with 6-OHDA has been reported to increase endogenous GABA levels (Tanaka et al. 1986; Lindefors et al. 1989), to increase potassium  $(K^+)$ -stimulated in vivo release of GABA (Lindefors et al. 1989; Tossman et al. 1986), and to increase the activity of glutamic acid decarboxylase (GAD), the enzyme responsible for converting glutamic acid to GABA (Nagy et al. 1978; Segovia et al. 1990; Samuel et al. 1988), and GAD mRNA levels (Vernier et al. 1988; Lindefors et al. 1990; Segovia et al. 1990; O'Connor et al. 1991). In neonatal rats, bilateral DA depletions have also resulted in increased tissue GABA levels in the striatum as well as in target regions of the striatopallidal and striatonigral projections, namely, the globus pallidus and SN 2-3 months after the lesion (Molina-Holgado et al. 1993a, b). This increase in GABA levels in the neonatally lesioned striatum was found to correspond to increased GAD mRNA expression (Soghomonian 1993; Laprade and Soghomonian 1999). Because nigrostriatal DA terminals that synapse on GABAergic mediumsized neurons exert an inhibitory effect over striatal GABA neurons (van der Heyden et al. 1980; Reid et al. 1990), attenuation of this inhibition after lesioning DA neurons is likely responsible for the increase in striatal GABA levels. In this respect, studies have shown that enhanced GABAergic activity after neonatal 6-OHDA lesions is influenced substantially by DA receptor status, since irreversible blockade of DA receptors by N-ethoxycarbonyl-2ethoxy-1,2-dihydroquinoline (EEDQ) decreased both D<sub>1</sub> and D<sub>2</sub> sites (73-84 %) in control and neonatal lesioned striatum, but increased GABA levels by 25 % only in rats that received 6-OHDA after birth (Molina-Holgado et al. 1995). Further, following acute inhibition of DA synthesis or catabolism, GABA levels remain unchanged. While not found in control rats, a positive correlation occurred between increased GABA levels and the density of  $D_2$  receptors, but not  $D_1$  receptors, in the striatum of neonatal

6-OHDA-lesioned rats, a finding suggestive of a direct role for  $D_2$  receptors in this regulation (Molina-Holgado et al. 1995). The increase in  $D_2$  receptors in the striatum, presumably on GABAergic neurons that project to the globus pallidus (striatopallidal pathway) may play a key role in maintaining this permanent increase in GABAergic transmission in rats with neonatal 6-OHDA lesions.

Altered GABA Receptors. Accompanying the 6-OHDA increase in GABAcontaining neurons is an increase in density of some GABA receptors in the striatum of neonatal lesioned rats (Molina-Holgado et al. 1995) – a change which likely produces plasticity of GABA-mediated signaling in neurons that form the major output systems of the striatum (i.e., the striatonigral and striatopallidal neurons). In rats lesioned unilaterally with 6-OHDA as adults, GABA<sub>A</sub> receptors were decreased in the striatum and globus pallidus, but increased in the SNr and entopeduncular nucleus (Pan et al. 1985). In neonatal 6-OHDA-lesioned rats,  $GABA_A$  receptors labeled with the radioligand [<sup>3</sup>H] muscimol were increased in the striatum, but not in the cerebral cortex or thalamus (Molina-Holgado et al. 1993a). It has been postulated that the increased  $[^{3}H]$  muscimol labeling in the striatum of neonatal lesioned rats targets GABAA receptors situated on hyperinnervated 5-HT fibers in the rostral striatum (Molina-Holgado et al. 1993a). Intracerebral microdialysis studies have shown that the striatal 5-HT hyperinnervation is not associated with enhanced basal levels of extracellular 5-HT (Jackson and Abercrombie 1992).

Changes in Other Amino Acid Transmitters. In addition to GABA, studies have demonstrated lasting increases in several other amino acids in the striatum and other regions of the basal ganglia in neonatal 6-OHDA-lesioned rats (Molina-Holgado et al. 1993a, b). In the striatum, aspartic acid and glycine were increased 3 months after the lesion, while in the globus pallidus, taurine, glutamine, aspartic acid, serine, and alanine were elevated (Molina-Holgado et al. 1993a, b). In the SN, only increases in taurine, glutamic acid, and glutamine were documented. When the amino acid profiles were compared to the relative amino acid composition of the tissues, the ratio of glutamic acid to GABA was decreased in the striatum and SN, but remained unchanged in the globus pallidus (Molina-Holgado et al. 1993b). Since the glutamic acid/GABA ratio can serve as an index for precursor product availability for GABA (Molina-Holgado et al. 1993b; Berl 1973), the decreased glutamic acid/GABA ratio implied an enhanced inhibitory influence relative to glutamic acid-mediated excitations in the striatum and SN (Molina-Holgado et al. 1993b). Such persistent and regionally distinct biochemical adjustments in amino acids within these basal ganglia regions demonstrate the extraordinary ability of the developing system to reorganize itself after early neonatal damage to DA neurons.

#### 4.5 Altered Cholinergic Function

Studies using the technique of in vivo microdialysis have demonstrated a decrease in the release of endogenous acetylcholine (ACh) in the striatum in response to  $D_2$  receptor agonists administered either systemically or locally through the dialysis

probe (Damsma et al. 1990; de Boer et al. 1990; De Boer et al. 1992). This finding is consistent with evidence that there is an important functional and anatomical relationship between  $D_2$  receptors and cholinergic interneurons in the striatum. In addition to changes in striatal  $D_2$  receptor density after 6-OHDA lesions to DA neurons, an increase in [<sup>3</sup>H]hemicholinium-3 binding, a marker for cholinergic terminal density has been reported (Joyce 1991b). Moreover, systemic or local administration of  $D_2$  blockers produces an increase in endogenous ACh release in the striatum (Damsma et al. 1991; De Boer et al. 1992). A study by Guyenet et al. revealed that there is a loss of tonic DA inhibitory control over the release of ACh after 6-OHDA treatment in adult rats (Guyenet et al. 1975). This effect was found to be reversed several weeks later, suggesting that the DA control is eventually reinstated in the lesioned animals (Guyenet et al. 1975; MacKenzie et al. 1989). This reinstatement has been suggested to be dependent upon increases in both the number of  $D_2$  receptors and the activity of the high-affinity choline transport system associated with cholinergic terminals (Joyce 1991b).

#### 4.6 Altered Expression of Neuropeptides

The enkephalin and tachykinin (substance P and neurokinin A) peptides have been shown to be present in high concentrations in the basal ganglia (Graybiel and Ragsdalev 1983). As mentioned earlier, two predominant subpopulations of GABAergic output neurons have been identified in the striatum that have distinct neuropeptide profiles. The striatopallidal pathway expresses enkephalin, whereas the striatonigral pathway expresses tachykinins and dynorphin (Gerfen and Young 1988; Gerfen et al. 1990, 1991; Gerfen 1992). Early ultrastructural studies showed that nigrostriatal DA neurons directly synapse on the enkephalin- and tachykinin-containing GABAergic neurons in rats (Kubota et al. 1986a, b; Kawai et al. 1987). This observation is consistent with the work demonstrating that these neuropeptide systems are regulated by nigrostriatal DA input (Bannon et al. 1987; Basura and Walker 2000; Campbell et al. 2001; Engber et al. 1986, 1987; Sivam and Krause 1990; Walker et al. 1991; Young et al. 1986; Obeso et al. 2000).

6-OHDA Treatments on Neuropeptides. In adult unilateral 6-OHDA-lesioned rats, immunohistochemical studies showed that striatal enkephalin is increased ipsilateral to the lesion (Voorn et al. 1987; Engber et al. 1991; Taylor et al. 1992; Cenci and Bjorklund 1993; Campbell and Bjorklund 1994; Thal et al. 1983), while substance P is decreased in the striatum (Voorn et al. 1987; Engber et al. 1991; Sivam 1989, 1991; Sivam et al. 1987). Electron microscopic analysis revealed that the enkephalin-immunoreactive synaptic bouton profiles in the striatum are increased by the lesion, without affecting the boutons on postsynaptic targets (Ingham et al. 1991). Furthermore, the unilateral 6-OHDA lesions were found to be accompanied by an increase in mRNA enkephalin levels (Gerfen et al. 1991; Campbell et al. 1992; Cenci et al. 1993; Nisenbaum et al. 1994a, b; Voorn et al. 1994; Zeng et al. 1995) and a decrease in preprotachykinin mRNA levels

(Campbell et al. 1992; Campbell and Bjorklund 1994; Nisenbaum et al. 1994a, b; Zeng et al. 1995). Studies also showed that transplants with DA-rich fetal cells are able to reverse the increased striatal enkephalin levels (Abrous et al. 1993b; Bal et al. 1993) consistent with striatal enkephalin levels being controlled by nigrostriatal DA neurons (Bannon et al. 1987; Basura and Walker 2000; Campbell et al. 2001; Engber et al. 1992; Gerfen et al. 1990; Hanson et al. 1981; Li et al. 1987; Sivam 1989; Sivam et al. 1986, 1987; Sivam and Krause 1990; Walker et al. 1991; Young et al. 1986; Obeso et al. 2000).

Although bilateral 6-OHDA lesions in both neonatal and adult rats have consistently increased both enkephalin protein and its mRNA levels (Sivam et al. 1986, 1987, 1991; Sivam and Krause 1990; Kurumaji et al. 1988; Cimino et al. 1991; Soghomonian 1994), only neonatal lesions consistently produce a decrease in the tachykinin peptides substance P and neurokinin A in the striatum (Sivam 1989; Sivam et al. 1987, 1991; Sivam and Krause 1990). Snyder-Keller and colleagues reported that the neonatal DA depletions do not affect the patchy distribution of substance P immunostaining, nor do they affect the distribution of enkephalin in the matrix of the striatum (Snyder-Keller 1991), suggesting that the basic organization of the striatal patch and matrix compartments develop normally in the absence of DA innervation early in development. In addition, while neonatal DA depletion was found to enhance the release of ACh, it has no effect on the action of substance P- or neurokinin A-induced ACh release in rat striatal slices (Perez-Navarro et al. 1993), indicating that striatal cholinergic neurons are modulated by tachykinin peptides independently of the presence of nigrostriatal DA neurons.

The effect of adult and neonatal 6-OHDA lesions on enkephalin, tachykinin, and other peptide levels has also been examined in the SNr, globus pallidus, and nucleus accumbens of rats. Immunohistochemistry and in situ hybridization studies showed that enkephalin and its mRNA levels are enhanced in the SNr after either adult unilateral or neonatal 6-OHDA lesions (Sivam et al. 1987; Vankova et al. 1991; Gerfen et al. 1991). In rats with adult unilateral lesions, neurotensin was reported to be increased and substance P reduced in the SNr (Taylor et al. 1992). In the SNr of rats with neonatal 6-OHDA lesions, no change in neurotensin was reported (Luthman et al. 1990a), although both substance P and neurokinin A were significantly reduced, an effect not observed in adult bilateral-lesioned rats (Sivam et al. 1987). Moreover, enkephalin and its mRNA (Schuller et al. 1999), neurotensin, and substance P were all increased in the globus pallidus of adult rats with unilateral lesions, but the increases were found within distinct neuronal subpopulations (Martorana et al. 2003). The contents of mRNA for enkephalin and peptides for leu- and met-enkephalin were increased, whereas mRNA for dynorphin and substance P were decreased in more rostral parts than caudal parts of the nucleus accumbens after adult unilateral 6-OHDA lesions (Taylor et al. 1992; Voorn et al. 1994). Similarly, neonatal 6-OHDA lesions produced reductions in substance P in the nucleus accumbens (Luthman et al. 1990a). Generally, the changes in peptide levels observed in the SNr, globus pallidus, and nucleus accumbens appear to reflect the peptide and mRNA changes documented in the striatum. A functional relationship between the nigrostriatal DA system and enkephalin or tachykinin

systems in the basal ganglia has been suggested by several investigators (Hong et al. 1978a, b, 1979, 1985; Joyce and Iversen 1979; Bannon et al. 1986; Hanson et al. 1981; Tang et al. 1983).

Evidence has also indicated the participation of corticostriatal afferents and cholinergic function in the regulation of these neuropeptides in the striatum. For example, frontal cortical transections after 6-OHDA treatment in adult animals were found to reverse the lesion-induced increase in preproenkephalin mRNA in the medial striatum, but have no effect on the upregulation found in the lateral striatum (Campbell and Bjorklund 1994). The decrease in preprotachykinin mRNA within the striatum was unaffected by interruption of the corticostriatal afferents (Campbell and Bjorklund 1994). Systemic administration of the muscarinic cholinergic receptor antagonist scopolamine to adult 6-OHDA-lesioned rats prevented the increase in enkephalin mRNA levels in the striatum (Nisenbaum et al. 1994b). Similarly, high doses of systemic scopolamine blocked the lesion-induced decrease in striatal substance P mRNA levels, possibly as a result from an action of scopolamine within other brain areas (Nisenbaum et al. 1994b). In addition to cortical influences, neuropeptide changes induced by adult unilateral 6-OHDA lesions have been shown to be inhibited by lesions of the subthalamic nucleus (Delfs et al. 1995), suggesting that this region can influence peptide function in the striatum through polysynaptic pathways. This latter conclusion is based upon the observation that lesions of the subthalamic nucleus inhibit neuropeptide changes induced by the 6-OHDA in brain regions that do not receive a direct input from the subthalamus.

In addition to the enkephalin and tachykinin peptides, 6-OHDA lesions have also been shown to alter levels of other peptides, including cholecystokinin, neurotensin, and somatostatin in the striatum. In adult rats, nigrostriatal DA depletion induced cholecystokinin and neurotensin mRNA in the striatum (Schiffmann and Vanderhaeghen 1992; Engber et al. 1992). The cholecystokinin mRNA expression was limited to dorsolateral parts of the ipsilateral denervated striatum (Schiffmann and Vanderhaeghen 1992). The increase in cholecystokinin was particularly unexpected, since cholecystokinin co-localizes with TH in the SNpc and VTA, even though not all DA neurons contain this peptide (Savasta et al. 1989). Furthermore, dynorphin mRNA expression in striatal patch neurons showed a marked reduction, while matrix neurons showed no change after lesioning of DA neurons in adult animals (Gerfen et al. 1991). Finally, a unilateral 6-OHDA lesion performed in neonatal rats had no effect on somatostatin mRNA (Cimino et al. 1991), but was found to increase neurokinin B, a member of the tachykinin family, on the ipsilateral side of the lesion (Burgunder 1991).

Drug Challenges on Neuropeptides After 6-OHDA. Treatment with DA agonists further implicated nigrostriatal DAergic control of neuropeptides located in the striatum. It has been shown that chronic treatment with either a  $D_1$  or  $D_2$  receptor agonist in adult unilateral 6-OHDA-lesioned rats affects several neuropeptides, but in distinct ways. Systemic administration of the  $D_2$  receptor agonist quinpirole inhibited the increase in enkephalin mRNA levels in these lesioned animals (Nisenbaum et al. 1994b; Engber et al. 1992). Moreover, systemic administration and direct infusion of quinpirole for 7 days to lesioned animals elevated induced increase in neurotensin and enkephalin (Nisenbaum et al. 1994b; Engber et al. 1992). Administration of the  $D_1$  receptor agonist SKF-38393 had no effect on enkephalin, but reduced somatostatin and neuropeptide Y levels, and increased neurotensin and dynorphin levels in the striatum of unilateral 6-OHDA-lesioned animals (Engber et al. 1992). These findings suggested that a reciprocal relationship exists between the function of these neuropeptides and the DA receptor subtypes in rats with a unilateral lesioning of DA neurons. In neonatal lesioned animals, administration of SKF-38393 beginning 24 h after 6-OHDA treatment reversed the lesion-induced increase in enkephalin and decrease in substance P in the striatum, while quinpirole (LY-17155) had no effect (Sivam and Cox 2006), suggesting an important role for  $D_1$  receptors in the postnatal development of enkephalin and substance P systems in this region. Kaakkola and colleagues found that circling behavior induced by an enkephalin analog, FK 33-824, microinjected into the SNr is antagonized by 6-OHDA lesions, suggesting that the response is mediated by the nigrostriatal DA pathway (Kaakkola 1980). In contrast, increased locomotor and rearing behavior in rats induced by microinjection of the peptidase-resistant enkephalin analog D-ala-met-enkephalamide (DALA) into the nucleus accumbens was found to be unaffected by the destruction of the mesolimbic DA system with 6-OHDA, indicating that this action occurred independent of an intact mesolimbic DA system (Kalivas et al. 1983).

#### 5 Behavioral Supersensitivity After 6-OHDA Lesions

#### 5.1 Hyperlocomotor and Stereotyped Responses to DA Agonists

Since the early 1970s, it has been established that all treatments resulting in the inhibition or loss of DA transmission in the adult striatum induce a behavioral supersensitivity to agonist administration. This phenomenon is most commonly examined by testing rats for their rotational response to mixed D<sub>1</sub>/D<sub>2</sub> receptor agonists (e.g., apomorphine) following adult unilateral lesions of the nigrostriatal DA pathway (Schwarting and Huston 1996). The behavioral supersensitivity after adult unilateral 6-OHDA lesions has been associated with the increase in  $D_2$ receptor number found in the striatum, suggesting that the change in  $D_2$  receptor density is one mechanism by which the DA denervation-induced supersensitivity occurs (Creese et al. 1977; Marshall and Ungerstedt 1977). Although rats unilaterally lesioned with 6-OHDA provide a useful model to examine drug actions in the CNS, one major problem is that the behavioral response to DA agonists cannot be examined in non-lesioned animals. An additional problem is that the intact control hemisphere may undergo a rapid reorganization (morphological and/or neurochemical) subsequent to the unilateral DA depletion which would allow for compensatory function on the contralateral side of the lesion (Tillerson et al. 2001; Woodlee et al. 2008), necessitating the assessment of biochemical changes in non-lesioned striatum as well in order to avoid a possible misinterpretation of data.

Bilateral destruction of DA neurons in adult rats has also been used to examine supersensitive responses to DA agonists. While systemic administration of low doses of apomorphine and L-dopa in controls produces a minor increase in locomotor responsiveness that is accompanied by sniffing, these  $D_1/D_2$  receptor agonists induce a major increase in locomotion in the bilaterally lesioned rats. The elevated locomotor response to apomorphine in the adult rats was found to be blocked by haloperidol, *cis*-flupentixol, and SCH-23390 (Breese et al. 1985a). Likewise, these adult bilaterally lesioned rats exhibited a major increase in activity when administered a  $D_2$  agonist, with only a variable, but modest limited increase being observed after  $D_1$  agonist administration (Breese et al. 1985a). In this latter case, haloperidol prevented the locomotor response to the  $D_2$  agonist, but SCH-23390 did not (Breese et al. 1985a).

Systemic treatment of intact adult rats with amphetamine or apomorphine also induces a repetitive occurrence of distinct stereotypic behaviors (Costall and Naylor 1974; Costall et al. 1975, 1977; Creese and Iversen 1974). The DA agonist-induced stereotypic behaviors are thought to reflect short-term changes in activity of DA receptors in the striatum of intact rats (Breese et al. 1984a; Costall and Naylor 1981; Silbergeld and Pfeiffer 1977). Increased locomotor activity and concomitant floor sniffing are the first components to occur following low doses of DA agonist (Breese et al. 1984a, 1985a), followed by repetitive limb and head movements such as rearing, paw treading, head bobbing, and floor/wall sniffing (Breese et al. 1984a, 1985a). At higher doses, locomotor components drop out and repetitive head and mouth movements (perioral responses) such as licking, biting, and gnawing predominate. Administration of apomorphine or an additional mixed  $D_1/D_2$  receptor agonist, L-dopa, to rats after adult bilateral 6-OHDA lesions produces a marked increase in locomotor activity over non-lesioned animals (Hollister et al. 1974) that is accompanied by stereotypic behaviors such as intense sniffing, rearing, paw treading, and head bobbing, reflecting a supersensitivity of DA receptors (Creese and Iversen 1974). In contrast, D-amphetamine does not produce any behavioral effects following its administration to these adult animals (Hollister et al. 1974). In unilaterally lesioned adults, d-amphetamine induces contralateral turning, a response opposite to that seen with apomorphine (Ungerstedt 1971b, d). Since amphetamine exerts its behavioral effects through the release of DA and blockade of its uptake into the nerve terminal, its ineffectiveness is the consequence of the 6-OHDA-induced DA loss (Breese et al. 1974).

Breese and colleagues were the first to show that certain behavioral responses to apomorphine and L-dopa in neonatal 6-OHDA-lesioned rats were significantly different from those in adult 6-OHDA-lesioned rats (Breese et al. 1984a, b, 1985b). While increased locomotion, sniffing, paw treading, and head nodding were behaviors characteristically associated with the mixed  $D_1/D_2$  agonist administration to rats bilaterally lesioned with 6-OHDA as adults or as neonates, selfbiting and a distinct behavior involving the repeated drawing of the forepaws up to and away from the nose referred to as "taffy pulling" were common unique responses in the neonatal lesioned rats. Both groups showed elevated rearing after either L-dopa or apomorphine treatment, indicative of their supersensitivity to these challenges (Breese et al. 1984a, b, 1985b). However, a high percentage of rats with neonatal 6-OHDA lesions exhibited self-injurious behavior (SIB) to a  $D_1/D_2$  DA agonist challenge, an effect not found in rats bilaterally lesioned with 6-OHDA as adults (Breese et al. 1984a, b. 1985b). While haloperidol reduced some behaviors including SIB induced by L-dopa in the neonatal lesioned rats, Breese and colleagues (1984a) further showed that *cis*-flupentixol was more potent than haloperidol in antagonizing the self-biting induced by L-dopa. In light of the report by Hyttel that *cis*-flupentixol was more selective for the  $D_1$  receptor site than the  $D_2$ receptor site (i.e., the haloperidol binding site) (Hyttel 1978), inhibition of SIB by *cis*-flupentized suggested that  $D_1$  receptors might be involved in the behavioral supersensitivity observed in rats depleted of DA during development (Breese et al. 1984a). Later studies showed that pretreatment with SCH-23390, a D<sub>1</sub> receptor antagonist, blocked self-biting and the SIB induced by L-dopa in neonatal 6-OHDA-lesioned animals, while D<sub>2</sub> receptor antagonists had a lesser effect (Breese et al. 1985a, 1989). Surprisingly, Breese et al. also reported that individual treatment with either a  $D_1$  or  $D_2$  receptor antagonist did not block the supersensitive locomotor response to L-dopa in adult or neonatal rats lesioned with 6-OHDA (Breese et al. 1985a). However, when these antagonists were combined, the enhanced locomotor activity induced by the L-dopa was virtually eliminated.

In subsequent studies, responses to specific  $D_1$  and  $D_2$  receptor agonists provided further evidence for distinct differences in responsive to these agonists depending on the age at which the DA was destroyed. When administered the specific  $D_2$ receptor agonist LY-171555 (quinpirole), adult rats with bilateral 6-OHDA lesions exhibited a much greater locomotor responsiveness than did neonatal lesioned animals (Breese et al. 1985a). In contrast, neonatal lesioned rats demonstrated a significantly greater behavioral responsiveness to administration of SKF-38393, a specific  $D_1$  receptor agonist, than did the adult lesioned group (Breese et al. 1985a). These findings suggested that the supersensitivity to DA agonists in the neonatal lesioned rats is primarily mediated by  $D_1$  receptors, whereas  $D_2$  receptors show greater sensitivity in the adult lesioned rats (Breese et al. 1985a). Such data provide further support for the view that the neonate and adult 6-OHDA lesions of the DA system induce adaptations that result in an altered functional relationship between  $D_1$  and  $D_2$  receptor function.

Altered Responses to DA Antagonists. As noted before, both neonatal and adult 6-OHDA-lesioned rats can be profoundly affected by doses of the catecholamine synthesis inhibitor  $\alpha$ -MT that have no impact in non-lesioned animals – a further demonstration of the importance of residual DA function in the lesioned animals (Cooper et al. 1973a; Luthman et al. 1995; Moy 1995; Potter and Bruno 1989; Rogers and Dunnett 1989a). Therefore, studies were undertaken to evaluate whether treatment with selective DA receptor antagonists to minimize the action of residual DA function in neonatal and adult lesioned rats would provide further support for the concept that age-dependent differences exist between the two groups. In this respect, adult rats lesioned as neonates show a subsensitivity to the D<sub>2</sub> receptor antagonist haloperidol and the D<sub>1</sub> receptor antagonist SCH-23390 compared to the action in adult lesioned rats (Bruno et al. 1984; Duncan et al. 1987). In non-lesioned controls and adult 6-OHDA-lesioned rats, Duncan and colleagues demonstrated that acute administration of either haloperidol or SCH-23390 produced a dose-dependent increase in akinesia (measured as movement on an inclined screen) (Duncan et al. 1987). In contrast, no significant alteration in immobility was observed in neonatal 6-OHDA-lesioned rats, even at a dose of SCH-23390 twice that necessary to produce akinesia in controls (Duncan et al. 1987). In another study, Bruno and colleagues reported subsensitivity to the akinesia-producing effect by haloperidol and fluphenazine, another neuroleptic, in neonatal rats lesioned at P3, but not those lesioned at P10, when tested during adulthood (Bruno et al. 1987). Rats with neonatal 6-OHDA lesions have also shown a subsensitivity to the effect of DA antagonist pimozide on lateral hypo-

thalamic self-stimulation (Stellar et al. 1988), a finding which led researchers

to suggest that a non-DA process mediates the observed behavioral sparing in these animals.

# 5.2 Priming of D<sub>1</sub> Agonist-Induced Supersensitivity

In rats lesioned as neonates, maximal supersensitivity of locomotor and stereotyped responses to SKF-38393 is achieved by administering multiple doses of the  $D_1$  agonist at weekly intervals over a period of 4 weeks (Breese et al. 1985a). This effect, referred to as "priming of  $D_1$  receptor supersensitivity," does not occur in rats with adult bilateral 6-OHDA lesions (Breese et al. 1985a). Such a sensitization process was not found after repeated  $D_2$  agonist treatment to neonatal 6-OHDA-lesioned rats, nor did a D<sub>2</sub> antagonist block the priming sensitization caused by SKF-38393 (Criswell et al. 1989). From studies involving intra-accumbens and intrastriatal antagonist-induced suppression of these behaviors, the enhanced locomotor effects could be attributed mainly to activation of D<sub>1</sub> receptors in the nucleus accumbens and stereotypies to activation of D<sub>1</sub> receptors in either the accumbens or the striatum (Breese et al. 1987a). Treatment with a full D<sub>1</sub> receptor agonist such as A-68390 or SKF-82958 can also induce priming in neonatal 6-OHDA-lesioned rats (Johnson et al. 1992; Bishop et al. 2003), although a major disadvantage is that large doses of these  $D_1$  receptor agonists desensitize the D<sub>1</sub> receptors. Still, at a reasonable dose, full D<sub>1</sub> agonists are more potent at inducing the priming phenomenon than SKF-38393 (Johnson et al. 1992; Bishop et al. 2003). The subsequent evaluation of priming of  $D_1$  receptor supersensitivity demonstrated that this effect was long-lasting and possibly permanent (Criswell et al. 1989). Another line of research focusing on increased oral activity in neonatal lesioned rats demonstrated a marked supersensitivity of these animals to SKF-38393 even at 8 months of age (Kostrzewa and Gong 1991). Furthermore, the repeated administration of a  $D_1$  receptor agonist to pre-weanling rat pups was found to cause a partial priming effect in adulthood (Gong et al. 1993a). Additionally, Kostrzewa and Kostrzewa found that guinpirole-induced vertical jumping induced during postnatal ontogeny was enhanced by neonatal 6-OHDA lesioning (Kostrzewa and Kostrzewa 2012).

Although adult animals given bilateral 6-OHDA lesions do not exhibit a priming effect with repeated DA agonist treatments, Morelli and DiChiara demonstrated that priming to SKF-38393 can be observed in adult animals given unilateral 6-OHDA lesions (Morelli and Di Chiara 1987). In this case, prior exposure to either a D<sub>1</sub> or D<sub>2</sub> receptor agonist results in an enhanced behavioral response to SKF-38393. While neonatal 6-OHDA-lesioned rats do not show a sensitization process to quinpirole after repeated treatments, they do, like adult unilateral lesioned animals, show a supersensitized response to SKF-38393 following a previous exposure to the selective D<sub>2</sub> agonist (Criswell et al. 1989). This evidence suggested that both neonatal and adult unilateral lesions may result in comparable adaptations of the DA system, while bilateral lesions in adult animals produce different compensatory changes.

#### 5.3 Functional Uncoupling of D<sub>1</sub> and D<sub>2</sub> Receptors by 6-OHDA

In non-lesioned rats, evidence has accumulated that  $D_1$  and  $D_2$  receptors are functionally "coupled." In this regard, Breese and colleagues reported that the locomotor stimulant effect of apomorphine, which has its primary effects on the  $D_1/D_2$  receptor sites, could be inhibited in non-lesioned animals by SCH-23390 (Breese and Mueller 1985; Breese et al. 1986), receptor antagonist with documented specificity for the  $D_1$  receptor (Billard et al. 1984; Molloy and Waddington 1984). Subsequently, Breese and Mueller demonstrated that this specific  $D_1$  receptor antagonist also prevented the locomotor response to the specific D<sub>2</sub> agonist, quinpirole, in control rats, just as did haloperidol, a D<sub>2</sub> receptor antagonist (Breese and Mueller 1985). In contrast to this finding, SCH-23390 was then found to be without effect on the response to this  $D_2$  agonist in rats with DA neurons lesioned with 6-OHDA. These findings suggested that the coupled functional interaction between  $D_1$  and  $D_2$  receptor subtypes to reach a maximal response appears to be dependent upon having an intact DA system (Breese and Mueller 1985). In additional studies, disruption of CAergic activity via the administration of reserpine and  $\alpha$ -MT was found to reduce the ability of SCH-23390 to block the effects of quinpirole in non-lesioned animals (Breese et al. 1986). These data collectively demonstrated clearly that  $D_1$  receptors can modulate  $D_2$  receptors in rats by a mechanism dependent upon intact DA function (Breese and Mueller 1985). Such disruption in the functional coupling of  $D_1$  and  $D_2$  receptors after impaired DAergic transmission has subsequently documented by several other investigators (Walters et al. 1987; Jackson and Hashizume 1987; Longoni et al. 1987).

In additional studies by Breese and colleagues in which the role of  $D_1$  and  $D_2$  receptor involvement of L-dopa induced SIB was explored,  $D_1$  and  $D_2$  agonists were administered alone and in combination to neonatal lesioned rats (Breese et al. 1985a). Repeated administration of the selective  $D_1$  agonist SKF-38393 to rats with neonatal 6-OHDA lesions at doses that produced priming did not lead to SIB, even though L-dopa-induced SIB could be blocked by a selective  $D_1$  receptor

antagonist. Moreover, treatment with the D<sub>2</sub> agonist quinpirole alone likewise did not lead to SIB (Breese et al. 1985a). Importantly when a low dose of quinpirole was combined with a low dose of SKF-38393, SIB was observed in the majority of neonatal 6-OHDA-lesioned rats treated (Breese et al. 1985a). This later finding suggested that while  $D_1$  receptor activation is essential for SIB to develop in rats with neonatal lesions, D<sub>2</sub> receptor activation, which also occurs with L-dopa treatment, is needed for the maximal expression of this SIB response. Johnson and Bruno showed that treatment with either a specific  $D_1$  or  $D_2$  receptor antagonist decreased food intake and induced akinesia in non-lesioned rats at doses that had no effect in neonatal lesioned animals (Johnson and Bruno 1992). Only when both agonists were administered together did the lesioned animals exhibit a response comparable to non-lesioned rats. A functional uncoupling of  $D_1$  and D<sub>2</sub> receptors by minimizing DA function may also explain the subsensitivity of neonatal 6-OHDA-lesioned rats to DA receptor antagonists (Johnson and Bruno 1992). Collectively, these observations provide further evidence 1990. for a functional relationship between the two DA receptor subtypes  $(D_1/D_2)$ coupling) being required for maximal effectiveness of DA function (Breese et al. 1985a).

The mechanisms involved in the  $D_1$  and  $D_2$  receptor interaction have been suggested to occur by receptor activation on discrete neurons (Gerfen et al. 1995; LaHoste et al. 2000) or by changes in D<sub>2</sub> receptor-mediated cholinergic interneuron activity with subsequent impact on  $D_1$  receptorexpressing striatonigral neuronal transmission (Di Chiara et al. 1994). These possibilities would seem consistent with the fact that GABAergic striatonigral and striatopallidal pathways have distinct DA receptor subtypes (Gerfen 1992) and that lesioning of DA neurons disrupting  $D_1/D_2$  receptor integration (Breese and Mueller 1985). However, more recent studies suggest that an interaction between  $D_1$  and  $D_2$  receptors occurring within the same cell may also be involved. For example, single medium spiny-to-medium spiny neuron synaptic connections in neonatal striatal cultures have been reported to exhibit modulation by both  $D_1$  and  $D_2$  receptor agonists (Geldwert et al. 2006), and the selective activation of  $D_1$  or  $D_2$  receptors in single striatal neurons has been shown to differentially modulate tetrodotoxin-sensitive sodium channels (Aizman et al. 2000). It is also now known that  $D_1$  and  $D_2$  receptors can form a heteromeric complex via a direct protein-protein interaction on medium spiny neurons that requires activation of both DA receptor subtypes for activation of the G-protein-coupled pathway (Rashid et al. 2007), an observation consistent with the striatal GABA ergic neurons that co-expresses both  $D_1$  and  $D_2$  receptors (Perreault et al. 2011). Consequently, possibly neurons with single as well as both DA receptor subtypes could account for the "uncoupling" of DA function described with lesioning. Regardless, the uncoupling of  $D_1$  and  $D_2$  receptor function has been suggested to be linked to a number of psychiatric illnesses related to DA dysfunction, including schizophrenia, drug abuse, and depression (Lee et al. 2009; Pei et al. 2010; Fuxe et al. 2008).

## 5.4 Hyperlocomotor Responses to NMDA Antagonists

Distinct pharmacological profiles also emerge when neonatal and adult 6-OHDAlesioned rats are administered MK-801, a noncompetitive antagonist of the N-methyl-D-aspartate (NMDA) receptor (Criswell et al. 1993; Morelli and Di Chiara 1990a). Rats lesioned as neonates show a latent supersensitivity to the hyperlocomotor stimulant action of MK-801 and CGS-19755, a competitive NMDA receptor antagonist. Rats lesioned as adults, either bilaterally or unilaterally, do not show this enhancement in MK-801-induced locomotor activity (Criswell et al. 1993; Morelli and Di Chiara 1990a; Carlsson and Carlsson 1989; Klockgether and Turski 1990; Morelli et al. 1992). Because pretreatment with  $\alpha$ -MT inhibits some, but not all, of the stimulant effects of MK-801 in neonatal lesioned rats (Criswell et al. 1993), a portion of the increased locomotor response can be attributed to some mechanism other than catecholamine or DA release. Additional evidence that the effects of MK-801 are mediated by at least two distinct mechanisms arises from the observation that the coadministration of  $D_1$  and  $D_2$ receptor antagonists attenuates some, but not all, of the actions of the NMDA antagonist in neonatal lesioned animals (Criswell et al. 1993). In contrast,  $\alpha$ -MT pretreatment in rats with adult lesions not only prevented the effects of MK-801 but also markedly attenuated any locomotor response (Criswell et al. 1993). The severity of  $\alpha$ -MT-induced akinesia seen in the adult lesioned rats further underscores the importance of the residual DAergic activity in these animals.

Studies have also shown that MK-801 and other NMDA antagonists can potentiate the effects of DA agonists. In non-lesioned rats pretreated with reserpine and  $\alpha$ -MT, the ability of L-dopa to diminish the resulting motor symptoms can be markedly potentiated by administering a single dose of MK-801 or CPP (a competitive NMDA antagonist), which on their own has no observed effects (Klockgether and Turski 1990). MK-801 also improves the effects of SKF-38393 on contralateral turning in rats given adult unilateral 6-OHDA lesions (Morelli and Di Chiara 1990a; Morelli et al. 1992), although the effects of MK-801 on L-dopainduced turning remain variable (Morelli et al. 1994; Spooren et al. 2000; Paquette et al. 2010; Dupre et al. 2008). Interestingly, MK-801 can prevent L-dopa-induced SIB in a dose-dependent manner in neonatal 6-OHDA-lesioned rats, while potentiating some stereotyped behaviors, i.e., paw treading, paw licking, and rearing, produced by DA agonists (Criswell et al. 1990). It has been postulated that the differential effects of MK-801 in DA-depleted rats are owed to distinct interactions between MK-801 and the  $D_1$  and  $D_2$  receptor subtypes (Morelli and Di Chiara 1990a, b; Morelli et al. 1992).

In line with these earlier findings with MK-801 and CGS-19755, more recent studies have revealed that neonatal 6-OHDA-lesioned rats also exhibit supersensitivity to the effects of phencyclidine (PCP), another NMDA receptor antagonist (Moy and Breese 2002). The PCP supersensitivity, displayed as an enhanced locomotor activity, was most pronounced after repeated PCP treatment or after a brief priming regimen with a  $D_1$  receptor agonist. This latter observation was

consistent with the process of sensitization of PCP that might be linked to changes in D<sub>1</sub> receptor function as noted in Sect. 5.5 to follow. Nonetheless, non-DAergic mechanisms may play a more important role in the acute attenuation of PCP effects in sensitized animals, since the enhanced locomotor responses in the neonatal lesioned animals could be blocked by the atypical antipsychotic agent olanzapine, a compound with activity at D<sub>1</sub>, D<sub>2</sub> and specific 5-HT receptors (Bymaster et al. 1996), but not by haloperidol or SCH-23390 (Moy and Breese 2002). Later studies showed that the selective 5-HT<sub>2A/2C</sub>-receptor antagonist ketanserin could significantly reduce PCP activation in lesioned rats, suggesting that the efficacy of olanzapine against PCP-induced sensitized responses may be mediated by one of these 5-HT<sub>2</sub> receptor subtypes (Moy et al. 2004).

# 5.5 NMDA Antagonist Inhibition of D<sub>1</sub> Agonist-Induced Supersensitivity

Apart from the stimulant effects on locomotor activity, blockade of the NMDA receptor can suppress some forms of neuronal plasticity, such as long-term potentiation (Morris et al. 1986), and can inhibit excitotoxic cell death induced by ischemia, trauma, and other insults (Hetman and Kharebava 2006). Important to this discussion of DA function, the long-lasting nature of priming of D<sub>1</sub> receptor supersensitivity to repeated doses of SKF-38393 in neonatal 6-OHDA-lesioned rats led to the hypothesis that this form of sensitization involves neural processes involving NMDA receptors similar to other forms of neuronal plasticity in CNS (Criswell et al. 1989). In line with this proposal, Criswell and colleagues showed that coadministration of MK-801 or CGS-19755 along with each priming dose of SKF-38393 blocked the behavioral sensitization (priming) observed by this repeated D<sub>1</sub> agonist administration to neonatal 6-OHDA-lesioned rats (Criswell et al. 1990). Since earlier work had demonstrated that the D<sub>1</sub> antagonist SCH-23390 also blocked the priming effect in neonatal lesioned rats (Criswell et al. 1989), it became clear that the concomitant activation of D<sub>1</sub> and NMDA receptors is likely necessary for priming of  $D_1$  receptor supersensitivity in neonatal lesioned rats to occur. The indication that NMDA receptor activation is also a necessary prerequisite for other responses that convey long-term consequences suggested that these phenomena share a common biochemical mechanism with D<sub>1</sub> receptor-mediated priming (see Sect. 5.2).

## 5.6 IEG and Signal Transduction Pathway Involvement in D<sub>1</sub> Agonist-Induced Supersensitivity

To date, a number of mechanisms potentially responsible for the development of priming of  $D_1$  receptor sensitivity in neonatal lesioned rats have been explored (Breese et al. 2005). The lack of any substantially significant changes in  $D_1$  receptor mRNA and/or expression in the striatum of rats after neonatal 6-OHDA lesions led

to the speculation that perhaps enhanced intraneuronal second messenger levels or efficiency likewise accompanied  $D_1$  agonist-induced priming in these animals (Gong et al. 1994). However, Gong and colleagues showed that basal as well as DA-, aluminum tetrafluoride-, and forskolin-stimulated adenylyl cyclase activities were not changed following repeated SKF-38393 treatment to neonatal lesioned rats during adulthood (Gong et al. 1994). Simson and colleagues also reported that SKF-38393-stimulated adenylyl cyclase activity was not altered in the striatum of these animals (Simson et al. 1992). These findings suggested that  $D_1$  receptorlinked adenylyl cyclase is not a major determinant of enhanced behavioral responses in neonatal lesioned rats to repeated  $D_1$  agonist administration. Likewise, other studies showed that adenosine 3–5'-monophosphate-regulated phosphoprotein (DARPP-32), a cyclic AMP-regulated phosphoprotein functionally linked to striatal  $D_1$  receptors (Barone et al. 1994), remained unaffected in the adult striatum of neonatal lesioned rats primed with multiple SKF-38393 treatments (Luthman et al. 1990a; Breese et al. 1994).

Because earlier studies had linked the proto-oncogene *c-fos* to behavioral supersensitivity to L-dopa or  $D_1$  agonists in rats either unilaterally or bilaterally lesioned with 6-OHDA as adults (Robertson et al. 1989a, b; Dragunow et al. 1991; Cole et al. 1993; Morelli et al. 1993), *c-fos* expression was examined in neonatal lesioned rats after  $D_1$  agonist treatment. Expression of *c-fos* occurred only after a dose of  $D_1$  agonist that was much higher than the behaviorally active dose and was not increased to a greater extent following repeated  $D_1$  agonist treatments (Johnson et al. 1992). Moreover, while NMDA receptor antagonists could block the  $D_1$  agonist-induced sensitization, it did not block *c-fos* expression in these animals (Johnson et al. 1992). Thus, it was concluded that *c-fos* is not a critical component in the priming phenomenon.

Mitogen-activated protein kinases (MAPKs) are intracellular mediators of signal transduction that are activated in response to a variety of extracellular stimuli (Johnson and Lapadat 2002; Pearson et al. 2001). There are three major classes of MAPKs: extracellular signal-regulated kinases (ERKs) (Sweatt 2001, p. 38; Johnson and Lapadat 2002), and stress-activated protein kinase/c-Jun amino-terminal kinase (SAPK/JNK) (Davis 2000). Among these, the most well studied are the closely related extracellular signal-regulated kinases 1 and 2 (ERK1/2). Biological processes involving ERK1/2 include stimulation of cell proliferation and survival, neoplastic transformation, neuronal differentiation, and plasticity (Seger and Krebs 1995; Paul et al. 1997). Several studies in both non-lesioned and adult 6-OHDAlesioned animals showed that the stimulation of  $D_1$  receptors can activate ERK1/2 to produce long-lasting adaptive changes in the brain (Vossler et al. 1997; Yao et al. 1998; York et al. 1998; Valjent et al. 2000; Gerfen et al. 2002). Thus, Papadeas and colleagues suggested that ERK1/2 may be involved in the adaptive processes that underlie the behavioral supersensitivity of neonatal lesioned rats to repeated  $D_1$ agonist treatments (Papadeas et al. 2004, 2008). Similar to reports in adult 6-OHDA-lesioned rats (Gerfen et al. 2002), the researchers found that single as well as multiple doses of SKF-38393 to rats with neonatal 6-OHDA lesions produces an acute, transient increase in ERK1/2 activity that peaks at 15 min following drug treatment (Papadeas et al. 2004). In conjunction with findings in adult lesioned animals, this finding suggested that the D<sub>1</sub> agonist-induced increases in locomotor activity could arise from strengthened ERK1/2-mediated signal transduction mechanisms involving in particular  $D_1$  receptor-expressing striatonigral neurons. The researchers also showed that the priming of  $D_1$  receptor sensitivity to multiple doses of SKF-38393 in rats with neonatal 6-OHDA lesions is coupled to a chronic, NMDA receptor-mediated ERK1/2 kinase activity (>7 days) in the medial prefrontal cortex and other cortical regions (i.e., the ventrolateral orbital, cingulate, motor, somatosensory, and piriform cortices) that promotes the phosphorylation of cAMP response element-binding protein (CREB), a transcription factor integral for neuronal plasticity (Lonze and Ginty 2002), and produces enduring structural changes in prefrontal cortical neurons (Papadeas et al. 2004, 2008). The lasting increase in ERK1/2 activity appears to be dependent upon  $D_1$  receptor function, as SCH-23390 antagonist pretreatment blocked the prolonged ERK1/2 response to repeated administration of SKF-38393 to neonatal lesioned rats (Papadeas et al. 2004). Importantly, the SCH-23390 did not inhibit ERK1/2 activation through its interaction with 5-HT<sub>2</sub> binding sites (Bischoff et al. 1988; McQuade et al. 1988), since systemic injections of the nonselective 5-HT<sub>2A/2C</sub> antagonist ketanserin prior to SKF-38393 had no effect on the prolonged ERK1/2 response.

In addition, intracerebroventricular or systemic pretreatment with the MAPK kinase (MEK)/ERK1/2 inhibitor SL327, and the structurally dissimilar PD 98059, prior to each of three of four weekly sensitizing doses of SKF-38393 was found to alter the character of the supersensitive behavioral response elicited by D<sub>1</sub> agonism (Papadeas 2006). Priming-induced locomotor behaviors such as running, walking, and trotting were augmented, while stereotyped behaviors such as rearing, paw treading, and taffy pulling were inhibited. It could be hypothesized that the augmentation of locomotor behaviors concomitantly inhibited the expression of rearing and paw-fixation behaviors in these animals (Bernardi et al. 1986; Chinen et al. 2006) or, alternatively, that the opposite effects of MEK inhibitor on locomotor and stereotyped behaviors may have occurred because the behaviors are driven by the activation of different neural systems. The microinjection of PD 98059 directly into the medial prefrontal cortex prior to SKF-38393 treatments inhibited the stereotyped behaviors, but did not affect locomotor behaviors (Papadeas 2006). This documentation suggested that forebrain ERK1/2 activation contributes to some but not all of the sensitized responses seen with D<sub>1</sub> agonist priming of neonatal lesioned rats and further that medial prefrontal cortical ERK1/2 activation may be critical for the sensitization of specific stereotyped behaviors exhibited by these animals.

## 5.7 Hyperlocomotor and Oral Responses to 5-HT Receptor Agonists

Evidence that neonatal 6-OHDA-lesioned rats show, in addition to the 5-HT hyperinnervation, increased 5-HT receptor density prompted several investigators to explore the role of 5-HT in spontaneous behavior and/or behavioral

responsiveness to DA agonists in these animals. Gong and Kostrzewa showed that systemic administration of the  $5-HT_{1B/2C}$  receptor agonist m-CPP could considerably increase oral activity in neonatal DA-depleted rats compared to control animals (Gong and Kostrzewa 1992). This enhanced oral response was dependent upon a selective supersensitivity of 5-HT<sub>2C</sub> receptors and was elevated to an even greater extent than that after D<sub>1</sub> agonist treatment in these animals (Gong and Kostrzewa 1992). In addition, the motor hyperactivity characteristic of these rats during adolescence (Shaywitz et al. 1976a; Erinoff et al. 1979; Heffner and Seiden 1982) was found to be reversed by systemic administration of the 5-HT<sub>2A</sub> receptor antagonists ketanserin and mianserin (Luthman et al. 1991), suggesting that the striatal increase in 5-HT<sub>2A</sub> receptors facilitates this abnormal behavior. Studies demonstrating the increased responses of striatal neurons to iontophoresed 5-HT and to 5-HT<sub>1B</sub> (m-CPP) and 5-HT<sub>2</sub> (DOI) agonists after neonatal 6-OHDA lesions further confirmed that increases in the corresponding 5-HT receptors were indeed functional (el Mansari et al. 1994). More recent studies have shown that the stimulation of 5-HT<sub>2</sub> receptors in neonatal or adult unilateral DA-depleted rats with the 5-HT<sub>2</sub> receptor agonist DOI can induce enhanced motor behaviors (Bishop et al. 2004; Bishop and Walker 2003), likely as a consequence of increased 5- $HT_{2A}$ receptor expression in the dorsal striatum (Radja et al. 1993a; Laprade et al. 1996; Basura and Walker 1999) and enhanced 5-HT<sub>2</sub> receptor-mediated activation of the direct striatal pathway (Gresch and Walker 1999; Basura and Walker 2000, 2001). Thus, in neonatal lesioned animals, striatal function may be enhanced significantly due to an elevated number of 5-HT receptors, even in the presence of extracellular 5-HT levels that remain near normal because of increased uptake (Jackson and Abercrombie 1992).

In the intact striatum, studies have consistently demonstrated that intrinsic 5-HT<sub>2</sub> receptors can modify DA function and have postulated divergent roles for 5-HT<sub>2A</sub> and 5-HT<sub>2C</sub> receptor subtypes (Lucas and Spampinato 2000; Porras et al. 2002). 5-HT<sub>2A</sub> antagonists reduce hyperlocomotion induced by cocaine, amphetamine, and 3,4 methylenedioxymethamphetamine (MDMA) (Kehne et al. 1996; O'Neill et al. 1999). Conversely, 5-HT<sub>2C</sub> receptor antagonists have been shown to enhance or reduce these effects depending upon which compounds and neural sites are studied (Filip and Cunningham 2002; Filip et al. 2004; Fletcher et al. 2002). While less is known about the interaction of these receptors following 6-OHDAinduced DA depletions, DA-related behaviors can be modified by 5-HT<sub>2</sub> receptor antagonists. For example, elevated oral dyskinesia induced by the ventral striatal infusion of SKF-38393 into neonatal 6-OHDA-lesioned rats was diminished by 5-HT<sub>2A/2C</sub> antagonism (Gong and Kostrzewa 1992; Plech et al. 1995). O'Boyle and colleagues demonstrated that hyperlocomotor activity induced by systemic or intrastriatal administration of SKF-82958 to adult unilateral 6-OHDA-lesioned rats (O'Boyle et al. 1989) can be reduced by pretreatment with the  $5HT_{2A/2C}$ antagonist ritanserin (Bishop et al. 2003). Further studies revealed that the  $D_1$ agonist-induced hyperlocomotor activity could be reduced by antagonism of striatal 5-HT<sub>2A</sub> receptors with M100907 (volinanserin), but not 5-HT<sub>2C</sub> receptors with RS102221 (Bishop et al. 2005). Volinanserin also significantly reduced

SKF-82958-induced hyperlocomotor activity in neonatal lesioned animals (Bishop et al. 2005). Taken together, these data suggest that adaptations in 5-HT receptors not only account for various aspects of spontaneous behavior, but can also influence supersensitive  $D_1$  receptors in rats with neonatal or adult unilateral 6-OHDA lesions.

## 5.8 Increased Stereotyped Responses to GABA Agonists

Differences in the behavioral profiles observed between neonatal and adult 6-OHDA-lesioned rats can also be seen following the administration of the GABA<sub>A</sub> receptor agonist, muscimol, into the SNr of these animals (Breese et al. 1987b). As mentioned earlier, the GABAergic neurons are projecting from the striatum into the SNr function as secondary mediators of DAergic activity (Albin et al. 1989; Gerfen et al. 1987). Additionally, DA neurons of the nigrostriatal pathway exert an inhibitory influence over these striatonigral neurons (Reid et al. 1990). Once the neurons of the nigrostriatal DA pathway are destroyed by 6-OHDA, the GABAergic activity in the SNr becomes enhanced. This presumably results in the increase in GABA levels observed in the SNr after neonatal 6-OHDA lesions (Molina-Holgado et al. 1993a). Elevated GABA activity in this region has been shown to play a role in SIB, as injections of muscimol into this region can elicit self-mutilation in normal animals (Baumeister and Frye 1986; Breese et al. 1987b). In both adult and neonatal DA-depleted rats, unilateral injections of muscimol into the SNr produced a supersensitive turning response as compared to non-lesioned animals (Breese et al. 1987b), a finding consistent with the reported increase in [<sup>3</sup>H]muscimol binding and immunostaining of  $GABA_A$  receptors in the SNr after adult unilateral 6-OHDA lesions (Pan et al. 1985; Katz et al. 2005). Only neonatal lesioned rats (and few non-lesioned rats) showed SIB and taffy-pulling behavior in response to bilateral injections of muscimol into the SNr (Breese et al. 1987b). Together, these data support the idea that lesions of DA neurons produce an increased functional responsiveness of  $GABA_A$  receptors in the SNr and that the increased susceptibility for SIB in neonatal DA-depleted rats is likely determined by neurons distal to the GABA<sub>A</sub> receptor complex in the SNr. Clearly, some other factor in the plasticity of these neurons to neonatal DA loss must account for the SIB that does not occur in the adult DA-depleted animals.

# 6 Modeling Specific Brain Disorders Using Rats with 6-OHDA Lesions

#### 6.1 Parkinson's Disease

The first of the clinical neurodegenerative disorders known to have reduced levels of DA and its metabolite HVA in the brain and CSF was PD (Hornykiewicz 1975).

PET studies with labeled agents that bind to DA uptake sites or to DA neurons via uptake sites such as fluorodopa confirmed that DA neurons were reduced (Eidelberg 1992; Shinotoh and Calne 1995). The availability of 6-OHDA has enabled the specific lesioning of DA neurons in adult rats to model the DA deficiency seen in PD (Ungerstedt 1968; Albin et al. 1989) and has allowed for the study of pathogenic mechanisms and pathophysiological aspects of PD as well as the testing of various neuroprotective compounds and innovative therapeutic strategies (Duty and Jenner 2011). Recently, adult lesioned rats have been used to investigate abnormal involuntary movements (AIMs) to prolonged use of L-dopa (Cenci et al. 1998; Winkler et al. 1999; Steece-Collier et al. 2003) and have provided invaluable information regarding the potential neurochemical and molecular mechanisms involved in L-dopa-related side effects in PD patients, such as dyskinesia (Cenci et al. 1998; Andersson et al. 1999; Sgambato-Faure et al. 2005; Pavon et al. 2006; Santini et al. 2007; Westin et al. 2007; Cenci and Konradi 2010). Further, it was revealed that striatal 5-HT terminals can cause or aggravate L-dopa-induced dyskinesias by

striatal 5-HT terminals can cause or aggravate L-dopa-induced dyskinesias by mishandling exogenous L-dopa and by releasing DA from the 5-HT terminals (Carta et al. 2007; Lindgren et al. 2010). In support of these actions related to 5-HT, administration of 5-HT<sub>1A</sub> and 5-HT<sub>1B</sub> agonists can have anti-dyskinetic effects (Carta et al. 2007; Lindgren et al. 2010). Currently, clinical investigations are underway to examine the efficacy of such 5-HT agonists in dyskinetic L-dopa-treated PD patients.

Still, the adult 6-OHDA-lesioned rat does not mimic all of the clinical features of PD. The depletion of DA, nigral DA cell loss, and neurobehavioral deficits have been successfully achieved with the model, but this lesioning does not appear to affect changes associated with other brain regions in PD, including olfactory structures, lower brain stem areas, or the locus coeruleus. Also, the adult 6-OHDA lesion does not produce or induce proteinaceous aggregates or Lewy-like inclusions like those seen in PD, although it has been reported that the 6-OHDA interacts with  $\alpha$ -synuclein (Blandini et al. 2008). Moreover, the nature of the symptoms in the unilateral adult 6-OHDA-lesioned rat, the most frequently used rodent model of PD, does not reflect that of the human disease. Despite these limitations, the adult 6-OHDA-lesioned rat model has contributed enormously to our understanding of PD and remains at the forefront of preclinical drug discovery for PD.

## 6.2 Lesch-Nyhan Syndrome

In addition to PD, the developmental disorder LNS (Lesch and Nyhan 1964; Nyhan 1968) is a second clinical disorder in humans with a documented destruction of central DA (Lloyd et al. 1981). The symptoms of LNS include motor dysfunction, mental impairment, and a particularly alarming form of SIB wherein patients bite away their own lips and fingers (Lesch and Nyhan 1964). The SIB is observed in about 85 % of LNS patients (Stout and Caskey 1989). The disease is caused by an almost complete deficiency of the enzyme hypoxanthine-guanine

phosphoribosyltransferase (HPRT) which also is associated with hyperuricemia. The main outcome of this genetically based defect is an increase in purine levels and a buildup of uric acid in the tissues. However, the severe neurological symptoms, including the behavioral abnormalities, are not affected by treatment that attenuates the overproduction of uric acid, indicating that some other consequence of the HPRT deficiency underlies the neurological deficits (Christie et al. 1982; Kopin 1981).

Research involving the examination of postmortem brain tissue revealed that LNS patients have a loss of 65–90 % of DA terminals in the mesolimbic and nigrostriatal pathways (Lloyd et al. 1981). Moreover, the spinal fluid of children with LNS has diminished levels of the DA metabolite HVA (Jankovic et al. 1988; Silverstein et al. 1985). Studies using PET technology to evaluate the integrity of DAergic neurons in LNS showed that a labeled marker for the DA transporter was markedly reduced in the brains of patients with LNS compared to either normal controls or patients with Rett's syndrome (Wong et al. 1996). In addition, labeled fluorodopa accumulation was reduced in the brains of patients with the disease (Ernst et al. 1996). It is likely, then, that these alterations in DA function account for the majority of the neurological symptoms observed in LNS (Jankovic et al. 1988; Lloyd et al. 1981).

Currently, the neonatal 6-OHDA-lesioned rat is the most widely accepted model to emulate the neonatal destruction of DA-containing neurons in LNS (Breese et al. 1984a, b, 1994), as these lesioned rats display characteristics that are similar to neurological changes and symptoms seen in patients with LNS. First, in both LNS patients and the neonatal lesioned rats, reduced DA during development is accompanied by an increase in striatal serotonin levels (Breese et al. 1984b; Lloyd et al. 1981; Stachowiak et al. 1984). Second, patients with LNS suffer from cognitive impairments (Lesch and Nyhan 1964), and considerable evidence suggests that neonatal lesioned rats do not learn behavioral tasks at the rate of non-lesioned animals (Archer et al. 1988; Smith et al. 1973). Third, similar to LNS patients, neonatal lesioned rats show an increased susceptibility for SIB (Breese et al. 1984b). Fourth, patients with LNS also show aggressive and combative behavior under the appropriate conditions (agitation and anxiety) (Lesch and Nyhan 1964), a characteristic feature of neonatal lesioned animals (Breese et al. 1984b). Fifth, patients with LNS are small in stature (Lesch and Nyhan 1964), as are neonatal lesioned rats compared to their non-lesioned counterparts (Smith et al. 1973). Sixth, antipsychotic drugs have little effect on SIB in LNS patients (Jankovic et al. 1988) and in neonatal lesioned rats administered sensitizing doses of L-dopa (Breese et al. 1984b). Finally, similar to neonatal lesioned rats, motor and behavioral symptoms are worsened when LNS patients are administered L-dopa (Jankovic et al. 1988). It is notable that the symptoms in PD patients are alleviated by L-dopa treatment, but neither PD patients nor adult 6-OHDA-lesioned animals exhibit self-biting or SIB. Collectively, the evidence clearly suggests that the early age of onset of loss of DA function is the key factor in the outcome of the DA deficiency in LNS (Breese et al. 1984b; Watts et al. 1982; Baumeister and Frye 1985).

#### 6.3 Other Psychiatric Conditions

There are three additional clinical disease states outlined below in which 6-OHDA lesions may be of potential usefulness in defining the basis of symptoms or sensitivity to drug responses observed after such lesions that are comparable to those observed in schizophrenia, drug addiction, or ADHD.

*Schizophrenia*. The enhanced susceptibility to NMDA antagonists production of psychosis observed in schizophrenia has led to the suggestion that a state of NMDA receptor hypofunction may be inherent to this psychiatric disorder (Javitt and Zukin 1991; Jentsch and Roth 1999; Olney 1989). Consequently, the exacerbated responsiveness to NMDA antagonists such as MK-801 and PCP in the neonatal 6-OHDA-lesioned rats could serve as a model of this NMDA dysfunction in schizophrenia. Rats with neonatal 6-OHDA lesions have been shown to have impaired habituation and alterations in startle responses and sensory gating (Moy et al. 1994; Schwarzkopf et al. 1992, 1996), as well as a marked susceptibility to the stimulant impact of pharmacological challenges and environmental stressors (Moy et al. 1994; Schallert et al. 1989). These characteristics are also similar to those observed in schizophrenia (Javitt and Zukin 1991; Jentsch and Roth 1999; Malhotra et al. 1997; Olney et al. 1999). Thus, the neonatal 6-OHDA-lesioned rat could be a valuable model to investigate the neural basis of the enhanced negative responsive to NMDA-receptor antagonists observed in schizophrenics.

Drug Addiction. The development of a behavioral sensitization occurs in nonlesioned rats following chronic, intermittent exposure to psychostimulant drugs such as cocaine, amphetamine, and methamphetamine (Segal and Mandell 1974; Nishikawa et al. 1983; Epstein and Altshuler 1978). Furthermore, increasing evidence suggests that this sensitization to psychostimulant drugs involves alterations in the sensitivity of postsynaptic  $D_1$  receptors (McCreary and Marsden 1993; Kuribara 1995a, b; Yoshida et al. 1995; Henry and White 1991). For example, pretreatment with the D<sub>1</sub> receptor antagonist SCH-23390 antagonizes the locomotor-activating effects of repeated cocaine and methamphetamine treatment and prevents the development of sensitization (McCreary and Marsden 1993; Kuribara 1995a, b; Yoshida et al. 1995). Pretreatment with a  $D_2$  receptor agonist, at least in the case of cocaine-induced sensitization, has no such effect on the sensitization (McCreary and Marsden 1993). Pretreatment with NMDA receptor antagonists such as MK-801 also blocks the sensitized response to psychostimulant drugs (Karler et al. 1989; Wolf 1998). These characteristics are similar to those related to priming of the behavioral responsiveness of neonatal 6-OHDA-lesioned rats to repeated doses of  $D_1$  agonist (Criswell et al. 1989). In this respect, both  $D_1$  and NMDA receptor antagonists block the "priming" of neonatal 6-OHDA-lesioned rats to repeated  $D_1$  agonist treatments (Criswell et al. 1990). Finally, both forms of sensitization are accompanied by adaptive responses that converge upon common cellular and molecular signal transduction pathways (e.g., ERK1/2 and CREB) that are associated with  $D_1$  and NMDA receptor activation and long-term plasticity (Fornai et al. 2009; Chao and Nestler 2004; Valjent et al. 2000; Girault et al. 2007; Papadeas et al. 2004). Given these similarities, neonatal lesioned rats primed with

 $D_1$  agonist could provide a model for investigating further the basis of the priming of  $D_1$  receptor sensitivity that occurs in normal rats to multiple doses of psychostimulant drugs. As neural adaptations that underlie the sensitization process in rats are thought to contribute to the development of the compulsive patterns of drug craving that characterizes drug addiction in humans (Robinson and Berridge 2008), investigations of the neural adaptations that develop in response to  $D_1$ priming in neonatal lesioned animals could help uncover  $D_1$  receptor-specific processes involved in human addictive behavior, such as drug craving to stress. One major advantage of this model is the persistence of the sensitized response to  $D_1$  agonists following priming.

ADHD. The neonatal 6-OHDA-lesioned rat has also been proposed to be a useful model for ADHD. Similar to ADHD patients, rat pups lesioned with 6-OHDA exhibit hyperactivity (Shaywitz et al. 1976b) and impaired learning in a spatial discrimination task (Archer et al. 1988) which improves after methylphenidate or amphetamine treatment (Davids et al. 2002, 2003). In addition, as in ADHD, the neonatal lesioned rats show an initial decrease in spontaneous motor behavior when placed in a novel environment, but after repeated testing, increased activity is observed, thus providing useful information regarding the mechanisms that cause hyperactivity (Luthman et al. 1989a). The hyperactivity in neonatal 6-OHDA-lesioned rats is accompanied by decreased striatal DAT and increased D<sub>4</sub> receptor expression, as well as altered 5-HT function (Luthman et al. 1989a; Zhang et al. 2001). While the hyperactivity is not altered by DAT inhibitors, it can be greatly reduced by  $D_4$  antagonists as well as inhibitors of NET and 5-HT transporters (Davids et al. 2002, 2003; Zhang et al. 2001, 2002a). These findings suggested that psychostimulants reduce the hyperactivity of neonatal lesioned rats by inhibiting NET and 5-HT transporters rather than DAT. Consistent with this conclusion, it has been demonstrated that administration of the NET inhibitor atomoxetine strongly antagonizes motor hyperactivity in juvenile rats with neonatal 6-OHDA lesions (Moran-Gates et al. 2005). When neonatally 6-OHDA-lesioned rats are lesioned in adulthood with 5,7-DHT, the level of hyperactivity is increased (Kostrzewa et al. 1994). Also, a 5-HT<sub>2</sub> receptor antagonist attenuates the spontaneous hyperlocomotor activity in neonatal 6-OHDA-lesioned rats (Brus et al. 2004). These findings further add support for involvement of the central 5-HT system in modulating hyperactivity. Furthermore, a possible role for 5-HT in stimulant treatment for ADHD has been proposed based on the demonstration of an inhibitory role of brain 5-HT systems in the actions of methylphenidate and amphetamine (Breese and Cooper 1975; Hollister et al. 1976).

### 7 Conclusion

The research on monoamine systems has greatly benefited from the discovery of 6-OHDA, the first compound implemented as a selective neurotoxin for CA-containing neurons (Porter et al. 1963; Ungerstedt 1968). Since the 1960s, 6-OHDA has been successfully employed in animals to biochemically dissect the roles of DA and NE neurotransmitter systems in the brain while at the same time

providing interesting models to study compensatory adaptations that occur in the metabolism, uptake mechanisms, sprouting, and plastic properties of the spared neurons and of the alternate neuronal projections that could replace the lesioned terminals. Eventual refinements in the use of 6-OHDA to selectively lesion DA neurons led to the proposal that DA-denervated rats provide a valuable model for human PD, a disease in which the loss of DA neurons is its main pathological hallmark. It soon became evident that while 6-OHDA-lesioned rats display many of the neurochemical characteristics found in human PD, they do not emulate all of the neurological features exhibited by patients. Clearly, the behavioral, neurochemical, morphological, and receptor alterations found in 6-OHDA-lesioned rats are dependent upon the degree of DA denervation, the site and route of 6-OHDA administration, and, most importantly of all, the age at which the lesioning is carried out. Furthermore, behavioral and receptor changes and the activation of associated signaling pathways are found not only in the central DA system as expected, but also in other transmitter systems, as demonstrated by some of the modifications in 5-HT and GABA and their receptors as reviewed above. Nonetheless, studies in both the adult and neonatal 6-OHDA-lesioned rat models still remain of interest, as they enable the examination of such alterations within a reasonable time frame, inevitably leading to the development and a better understanding of innovative therapeutic strategies to treat PD and other DA deficiency states like LNS, ADHD, and schizophrenia.

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# Dopamine and L-dopa as Selective Endogenous Neurotoxins

# Juan Segura-Aguilar, Ulises Ahumada-Castro, and Irmgard Paris

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#### Abstract

Selective neurotoxins have the ability to exert their neurotoxic effects in some specific neuronal systems. In dopaminergic neurons, the selectivity of exogenous neurotoxins depends on their affinity to the dopamine transporter. However, dopamine and 3,4-L-dihydroxyphenylalanine (L-dopa) are synthesized in dopaminergic neurons and are likewise able to induce neurotoxicity. The possible molecular mechanisms involved in dopamine and L-dopa neurotoxicity in dopaminergic neurons are discussed. Dopamine seems to be neurotoxic in dopaminergic neurons by undergoing oxidation to aminochrome, which is the precursor to neuromelanin. However,

e-mail: jsegura@med.uchile.cl

I. Paris

Department of Basic Sciences, Santo Tomas University, Viña del Mar, Chile

J. Segura-Aguilar (🖂) • U. Ahumada-Castro

Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

aminochrome can be neurotoxic when it forms adducts with proteins such as alphasynuclein, parkin, mitochondrial complexes I and III, actin, tubulin, and the dopamine transporter, or when aminochrome is one-electron reduced by flavoenzymes that use NADH, generating redox cycling with the concomitant depletion of energy and the formation of reactive oxygen species. L-dopa is also neurotoxic in cell cultures after oxidizing to a quinone species, but L-dopa seems to be a transient precursor of dopamine in that it is not able to induce neurotoxicity in vivo due to the efficient decarboxylation to dopamine catalyzed by amino acid decarboxylase. In fact, the only metabolite found in vivo is L-3-o-methyldopa, as detected in microdialysis experiments in animals treated with L-dopa. L-dopa is used in Parkinson's disease treatment, and it is still questionable whether L-dopa accelerates the degeneration of remaining dopaminergic neurons. It seems that L-dopa itself does not accelerate dopaminergic neuron degeneration because L-dopa is efficiently converted to dopamine, both in the peripheral and the central nervous systems. However, L-dopa induces dyskinesias in approximately 40 % patients with 4-6 years of treatment, and although the mechanism for L-dopa-induced dyskinesias is very complex, the rapid oscillation of striatal dopamine during L-dopa treatment has been found to be required for the induction of dyskinesias. The remaining dopaminergic neurons convert L-dopa to dopamine and release dopamine to the striatum under regulated conditions, but the majority of dopamine release to the striatum is mediated by serotonergic neurons without regulation, resulting in dyskinesias.

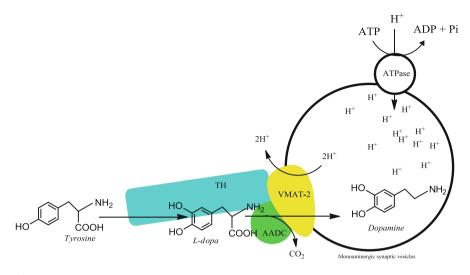
#### **Keywords**

Aminochrome • Dopamine • Dyskinesia • L-dopa • Metabolism • Neurotoxicity • Neurotoxins • Orthoquinones • Oxidation • Parkinson's disease

List of Abbre	viations
AADC	Aromatic amino acid decarboxylase
COMT	Catechol ortho-methyltransferase
DA	Dopamine
GST M2-2	Glutathione S-transferase M2-2
L-dopa	L-dihydroxyphenylalanine
MAO	Monoamine oxidases
TH	Tyrosine hydroxylase
VMAT-2	Vesicular monoaminergic transporter-2

## 1 Synthesis of L-dopa and Dopamine

De novo synthesis of dopamine is mediated by two enzymes using the amino acid tyrosine as precursor. The first step is catalyzed by the enzyme tyrosine hydroxylase (TH), in which tyrosine is used to form L-dihydroxyphenylalanine (L-dopa). L-dopa is decarboxylated by the enzyme aromatic amino acid decarboxylase (AADC) to generate dopamine and CO<sub>2</sub>. This dopamine is stored in monoaminergic

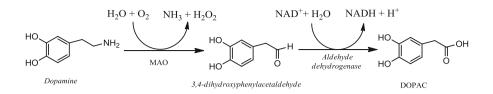


**Fig. 1** *Dopamine synthesis.* Dopamine synthesis from tyrosine is catalyzed by tyrosine hydroxylase (TH) and aromatic amino acid decarboxylase (AADC), and uptake into monoaminergic synaptic vesicles is mediated by VMAT-2

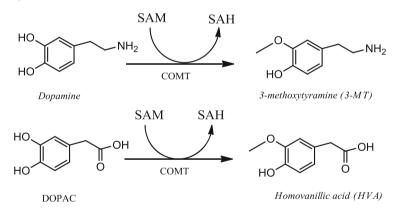
vesicles with a weak acid pH that prevents dopamine oxidation to o-quinone species, as dopamine autoxidizes at physiological pH due to the dissociation of the protons of hydroxyl groups. Dopamine oxidation to o-quinones after the synthesis of L-dopa and dopamine is prevented by the association of the enzymes TH and AADC with the vesicular monoaminergic transporter-2 (VMAT-2), which is present in the membranes of monoaminergic synaptic vesicles, generating a kind of complex that prevents free cytosolic dopamine (Cartier et al. 2010). Dopamine inside monoaminergic synaptic vesicles is hard protonated, preventing the oxidation of catechol groups to o-quinone. Monoaminergic synaptic vesicles express a vesicular ATPase that hydrolyzes ATP to ADP plus inorganic phosphate with one proton (H<sup>+</sup>) translocation into the vesicles, creating a proton gradient with a weak acid pH inside the vesicles (Guillot and Miller 2009; Fig. 1).

#### 1.1 Dopamine Degradation

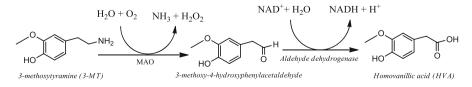
Dopamine accumulates inside monoaminergic synaptic vesicles for neurotransmission, which is an essential process in the control of movements. However, the catechol group of free cytosolic dopamine can be oxidized to neurotoxic *o*-quinones (Arriagada et al. 2004; Paris et al. 2010, 2011). To prevent this effect, dopamine participates in a neurotoxic reaction, and there are two types of enzymes that degrade dopamine. The first is the monoamine oxidases (MAO, E.C. 1.4.3.4), which catalyze the oxidative deamination of the dopamine amino group to 3,4-dihydroxyphenylacetaldehyde with concomitant formation of ammonia and hydrogen peroxide. Aldehyde dehydrogenase then catalyzes the oxidation of 3,4-dihydroxyphenylacetaldehyde to 3,4-dihydroxyphenylacetic acid (DOPAC) with the formation of a molecule of NADH.



The MAO enzymes have multiple isoforms (A and B forms) that are 70 % identical and are localized to the outer membranes of the mitochondria in neurons, glial cells, and other cells (Weyler et al. 1990; Shih et al. 1997). MAO-B is found in histaminergic and serotonergic neurons, as well as in astrocytes, while MAO-A is mainly found in catecholaminergic neurons (Westlund et al. 1988; Saura et al. 1994). MAO-A uses the compounds dopamine, noradrenaline, adrenaline, and serotonin as substrates, while MAO-B uses compounds such as phenylethylamine and tyramine (Strolin-Benedetti et al. 1992). The second enzyme that degrades dopamine is catechol ortho-methyltransferase (COMT; EC 2.1.1.6), which catalyzes the methylation of dopamine by using S-adenosylmethionine (SAM) as a cofactor to form 3-methoxytyramine and S-adenosylhomocysteine (SAH). COMT also plays a role in MAO-catalyzed degradation of dopamine, as DOPAC is methylated to homovanillic acid (HVA).



MAO also participates in dopamine degradation catalyzed by COMT because 3-methoxytyramine is a substrate for MAO, which catalyzes the oxidative deamination of 3-methoxytyramine to 3-methoxy-4-hydroxyphenylacetaldehyde. Aldehyde dehydrogenase catalyzes the oxidation of 3-methoxy-4-hydroxyphenylacetaldehyde to homovanillic acid with the concomitant formation of NADH.



COMT is expressed in pyramidal neurons, striatal spiny neurons, cerebellar Purkinje, and granular cells (Myöhänen et al. 2010), and it has two isoforms – a soluble form (S-COMT) and a membrane-bound form (MB-COMT). Inhibition of COMT by entacapone has been used in the treatment of Parkinson's disease, prolonging the half-life of L-dopa (Marin and Obeso 2010).

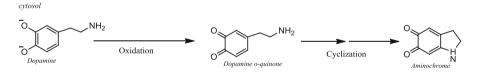
#### 2 Dopamine Neurotoxicity

Dopamine exposure at different cell cultures or cell lines resulted in the induction of cell death, suggesting a neurotoxic role for dopamine (Hoyt et al. 1997; Jeon et al. 2010). Dopamine-dependent cell death has been reported to be dependent on an apoptotic cell death mechanism (Ziv et al. 2001). Dopamine exposure of human neuroblastoma NMB cells, which are able to take up dopamine, induces morphological changes, such as cell shrinkage, apoptosis-like atrophy, accumulation of apoptotic particles, DNA fragmentation, and protein synthesis-dependent cell death (Simantov et al. 1996). A study conducted with SH-SY5Y neuroblastoma cells displayed caspase-9 and caspase-3 activation, cleavage of poly(ADP-ribose) polymerase, and nuclear condensation, accompanied by the activation of p38 mitogen-activated protein kinase. The presence of the antioxidant N-acetyl-Lcysteine prevents dopamine-induced p38 kinase activation, caspase-9 and caspase-3 cleavage, and subsequent apoptosis (Junn and Mouradian 2001). Overexpression of Bcl-2, which blocks physiological apoptosis, in PC-12 cells results in a marked resistance to dopamine-induced cell death (Offen et al. 1997). The activation of the JNK pathway precedes dopamine-induced apoptosis and is persistently sustained during the process of apoptosis. Overexpression of a dominant negative mutant SEK1, an upstream kinase of JNK, inhibits both dopamine-induced JNK activation and apoptosis (Luo et al. 1998). Another study reported that dopamine activates SAPK/JNK and p38, but not MEK or ERK/MAPK (Gómez-Santos et al. 2003). It has been reported that dopamine oxidation activates the DNA-binding activity of NF-kappaB and the suppression of NF-kappaB transcriptional activity in PC-12 cells, inducing apoptotic cell death during dopamine oxidation (Lee et al. 2001b). The formation of adducts between alpha-synuclein and the presynaptic human dopamine transporter facilitates the membrane clustering of the dopamine transporter (DAT), thereby accelerating cellular dopamine uptake and dopamine-induced cellular apoptosis (Lee et al. 2001a). Interestingly, parkin prevents dopamine-induced alpha-synuclein-dependent cell death by blocking DAT-mediated dopamine uptake, which is accelerated by alpha-synuclein (Moszczynska et al. 2007). Another study showed that parkin protects against dopamine toxicity by decreasing oxidative stress and the subsequent activation of the JNK/caspase pathway (Jiang et al. 2004). The apoptosis induced by dopamine in PC-12 cells was found to be accompanied by an impairment of mitochondrial bioenergetic functions (Jana et al. 2011). The treatment of human SH-SY5Y cells with dopamine induced an early increase in the expression of hypoxia-inducible factor-1alpha (HIF-1alpha) followed by increases in p53, Puma, and Bnip3, in which caspase-3, caspase-7, and PARP were activated after 12 h (Giménez-Xavier et al. 2009). The agonist of protein kinase A, forskolin, stimulated dopamine uptake in SK-N-SH cells and blocked dopamine-induced apoptosis (Liu et al. 2001). Dopamine-induced cell death was found to be preceded by a decrease in proteasome activity (Keller et al. 2000). Antioxidants were found to protect against dopamine-induced cell death (Junn and Mouradian 2001; Jana et al. 2011), and the glutathione-depleting compound L-buthionine sulfoximine enhanced dopamine-induced cell death (Stokes et al. 2000).

Why does dopamine induce neurotoxicity when it is an essential neurotransmitter for the control of movement? A possible explanation is that dopamine inside monoaminergic synaptic vesicles is completely inert because the protons of dopamine groups are hardbound to the oxygen as a consequence of the weak acid pH inside the vesicles, which is 2–2.4 pH units lower than the pH in the cytosol (Guillot and Miller 2009). However, the protons of dopamine hydroxyl groups are dissociated at cytosolic pH and can oxidize in the presence of oxygen, even in the absence of metal catalysis (Linert et al. 1996). The dopamine catechol structure oxidizes to orthoquinones, which can induce neurotoxicity.

#### 3 Dopamine Oxidation to Orthoquinones

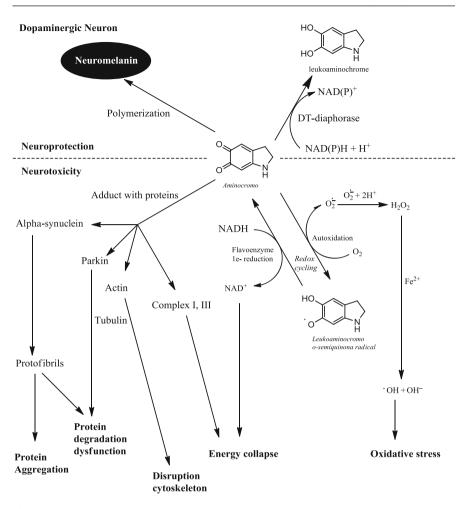
The protons of dopamine hydroxyl groups are dissociated when dopamine is in the cytosol at physiological pH and spontaneously oxidizes in the presence of oxygen and in the absence of metal traces (Linert et al. 1996). Dopamine oxidizes to dopamine *o*-quinone, which cyclizes to form aminochrome at physiological pH. Dopamine *o*-quinone is not stable at physiological pH because the amino group of dopamine *o*-quinone spontaneously rearranges and undergoes cyclization; thus, dopamine *o*-quinone is only stable at a pH below 2.0 (Segura-Aguilar and Lind 1989).



Dopamine oxidation can also be catalyzed by metals such as manganese(III), copper sulfate(II), iron chloride(III), and sodium periodate (Segura-Aguilar and Lind 1989; Paris et al. 2001, 2005a; Graham et al. 1978) or by enzymes such as prostaglandin H synthase, cytochrome P450, xanthine oxidase, lactoperoxidase, tyrosinase, and dopamine  $\beta$ -monooxygenase (Galzigna et al. 2000; Thompson et al. 2000; Segura-Aguilar 1996; Foppoli et al. 1997; Hastings 1995; Segura-Aguilar et al. 1998; Jimenez et al. 1984). It is important to remember that aminochrome formation is dependent on the presence of free cytosolic dopamine that can oxidize to aminochrome, and VMAT-2 and MAO prevent dopamine oxidation, as VMAT-2 mediates dopamine accumulation in monoaminergic synaptic vesicles and MAO catalyzes the degradation of dopamine.

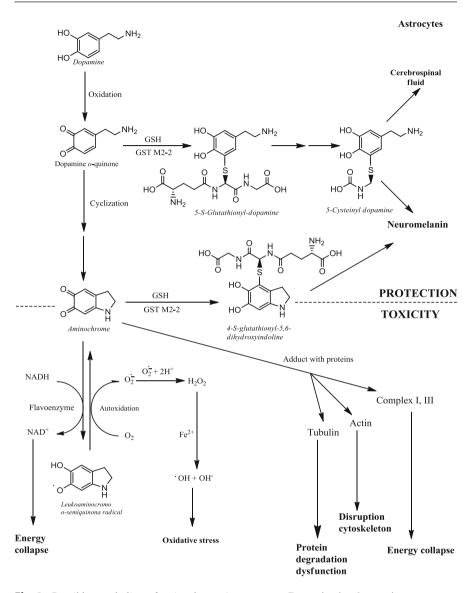
#### 4 Aminochrome Metabolism

Aminochrome is able to participate in five different reactions, including (i) the polymerization of aminochrome to neuromelanin. Neuromelanin is a pigment present in dopaminergic neurons, which is localized in the substantia nigra and accumulates with age (Zecca et al. 2002) in healthy individuals; it is also dramatically decreased in patients with Parkinson's disease. Therefore, neuromelanin seems to play a protective role in dopaminergic neurons because it chelates metals and binds proteins such as alpha-synuclein (Gerlach et al. 2003; Hong and Simon 2007; Fasano et al. 2006). The localization of neuromelanin in double-membrane vesicles seems to be very important for its protective role, as free neuromelanin has been found to be neurotoxic in cell cultures (Naoi et al. 2008) and extracellular neuromelanin induces microglial activation in the substantia nigra (Zhang et al. 2011: Fig. 2). (ii) Aminochrome also forms adducts with proteins. Aminochrome induces and stabilizes the formation of neurotoxic protofibrils of alpha-synuclein (Conway et al. 2001; Norris et al. 2005). Aminochrome forms adducts with other proteins, such as actin and  $\alpha$ - and  $\beta$ -tubulin, and it disrupts the architecture of the cytoskeleton and complexes I and III of the mitochondria (Paris et al. 2010; Van Laar et al. 2009). Dopamine o-quinone, the transient precursor of aminochrome at physiological pH, forms adducts with and inactivates parkin, which is a ubiquitin ligase of the proteasomal system (LaVoie et al. 2005); tyrosine hydroxylase; the human dopamine transporter; and tryptophan hydroxylase (Xu et al. 1998; Whitehead et al. 2001; Kuhn and Arthur 1998). (iii) Aminochrome can be one-electron reduced, catalyzed by flavoenzymes that transfer one electron by using NADH or NADPH ions. The leukoaminochrome o-semiquinone radical is extremely reactive under aerobic conditions, generating a redox cycle between the leukoaminochrome o-semiquinone radical and aminochrome (Baez et al. 1995; Segura-Aguilar et al. 1998). This redox cycling depletes the NADH and O<sub>2</sub> required for ATP production in the mitochondria or the NADPH required for the reduction of oxidized glutathione, which is an important antioxidant. There is significant evidence that supports the neurotoxic role of the one-electron reduction of aminochrome (Paris et al. 2001, 2005a, b, 2009, 2010, 2011; Arriagada et al. 2004; Fuentes et al. 2007; Díaz-Véliz et al. 2008; Muñoz et al. 2012a, b; Fig. 2). (iv) Aminochrome can be two-electron reduced to leukoaminochrome, catalyzed by DT-diaphorase (EC.1.6.99.2), which is a flavoenzyme that uses both NADH and NADPH as electron donors. DTdiaphorase has been proposed to play a protective role in aminochrome metabolism because this enzyme prevents the neurotoxic reactions of aminochrome, including the one-electron reduction of aminochrome and the formation of adducts with proteins (Arriagada et al. 2004; Lozano et al. 2010; Paris et al. 2011; Muñoz et al. 2012a, b; Segura-Aguilar et al. 2006; Cardenas et al. 2008; Paris et al. 2010; Fig. 2). (v) Aminochrome can be glutathione-conjugated by glutathione S-transferase M2-2 (GST M2-2) to 4-S-glutathionyl-5,6-dihydroxyindoline, which is a stable molecule that is resistant to biological oxidizing agents (Segura-Aguilar, et al. 1997; Baez et al. 1997). GST M2-2 also conjugates the precursor of aminochrome dopamine o-quinone to 5-glutathionyl-dopamine, preventing the



**Fig. 2** *Possible metabolism of aminochrome in dopaminergic neurons.* Aminochrome is able to participate in two neurotoxic reactions, such as the formation of aminochrome adducts with alpha-synuclein, parkin, actin, tubulin, and mitochondrial complexes I and III, along with aminochrome one-electron reduction to the leukoaminochrome o-semiquinone radical. However, aminochrome can polymerize into neuromelanin, or it can be two-electron reduced to leukoaminochrome, catalyzed by DT-diaphorase, a reaction that prevents aminochrome-induced neurotoxicity

formation of aminochrome (Dagnino-Subiabre et al. 2000). The 5-glutathionyldopamine is finally converted to 5-cysteinyl dopamine (Shen et al. 1996), which has been detected in the cerebrospinal fluid of Parkinson's disease patients and in dopamine-rich brain regions such as the caudate nucleus, putamen, globus pallidus, and substantia nigra, as well as in neuromelanin (Cheng et al. 1996; Rosengren et al. 1985; Carstam et al. 1991). Therefore, the conjugation of glutathione has been proposed to be a protective reaction against aminochrome neurotoxicity in astrocytes (Fig. 3).



**Fig. 3** *Possible metabolism of aminochrome in astrocytes.* Dopamine is taken up into astrocytes and is able to oxidize into dopamine *o*-quinone. GST M2-2 is able to conjugate this compound with glutathione to form 5-glutathionyl-dopamine, which is degraded to 5-S-cysteinyl dopamine, a compound that is found in the cerebrospinal fluid and in neuromelanin. Dopamine o-quinone cyclizes to aminochrome, which can also be conjugated with GSH by GST M2-2 to 4-S-glutathionyl-5,6-dihydroxyindoline, which is resistant to biological oxidizing agents. Both conjugations prevent aminochrome-induced neurotoxicity

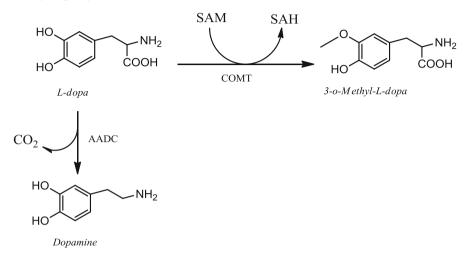
## 5 Aminochrome and Parkinson's Disease

The progressive loss of nigral dopaminergic neurons containing neuromelanin in the substantia nigra most likely begins long before the symptomatic phases of Parkinson's disease are present (Braak et al. 2004). Intensive research has been conducted over the past few decades to understand the molecular mechanism of the neurodegeneration of neuromelanin-containing dopaminergic neurons, with the discovery of mutations associated with Parkinson's disease. It is generally accepted that the mechanism that results in the loss of dopaminergic neurons in the substantia nigra involves the aggregation of alpha-synuclein to neurotoxic protofibrils, a dysfunction of protein degradation, mitochondrial dysfunction, oxidative stress, and neuroinflammation (Schapira 2011; Conway et al. 2001; McNaught et al. 2004; Cuervo et al. 2010; Schapira and Jenner 2011). There is growing evidence that supports the role of aminochrome in the degeneration of dopaminergic neurons that contain neuromelanin, including the following findings: (i) Aminochrome forms adducts and inactivates complexes I and III and isocitrate dehydrogenase, an enzyme of the citric acid cycle in the mitochondria (Van Laar et al. 2009). One-electron reduction of aminochrome depletes NADH, thereby decreasing ATP production in cell culture (Muñoz et al. 2012b). (ii) Aminochrome was found to induce the formation and stabilization of alpha-synuclein protofibrils (Norris et al. 2005). (iii) Aminochrome inactivates the proteasomal system of degradation of proteins by forming adducts with parkin (Zafar et al. 2006; La Voie et al. 2005). Interestingly, alpha-synuclein protofibrils inhibit chaperone-mediated autophagy (Xilouri et al. 2009) and the 26S proteasomal system (Zhang et al. 2008). Aminochrome forms adducts with tubulin (Van Laar et al. 2009), and one-electron reduction of aminochrome induces the aggregation of  $\alpha$ - and  $\beta$ -tubulin, disrupting the cytoskeletal structure (Paris et al. 2010; Muñoz et al. 2012a). Tubulin aggregation prevents the microtubule formation required for the fusion of autophagocytic vacuoles and lysosomes (Monastyrska et al. 2009). (iv) Aminochrome one-electron reduction induces oxidative stress (Arriagada et al. 2004). Interestingly, aminochrome is involved in four of the five mechanisms that are generally accepted to be involved in the degeneration of neuromelanincontaining dopaminergic neurons in Parkinson's disease.

## 6 L-dopa Metabolism

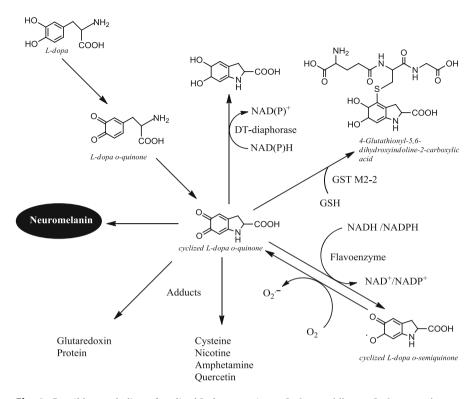
The amino acid tyrosine is used as a substrate in the production of L-dopa, a reaction catalyzed by tyrosine hydroxylase in dopaminergic neurons. However, L-dopa is immediately converted to dopamine in a reaction catalyzed by AADC. It has been reported that the enzymes TH and AADC are associated with the membrane-bound VMAT-2, forming a kind of complex to prevent L-dopa and dopamine from remaining free in the cytosol (Cartier et al. 2010). It is unknown whether free L-dopa in the cytosol is able to undergo metabolism under

normal conditions because in Parkinson's patients, L-dopa undergoes metabolism, an important part of treatment. L-dopa can be decarboxylated to dopamine even outside the central nervous system, and L-dopa can be converted to L-3-*o*-methyldopa by COMT (Okada et al. 2011).



## 7 L-dopa Neurotoxicity In Vitro

Experiments in vitro with cultured neurons exposed to L-dopa resulted in apoptosis that included cell shrinkage, membrane blebbing, and nuclear and DNA fragmentation (Melamed et al. 1998; Walkinshaw and Waters 1995). L-dopa was found to be neurotoxic when it increased the expression of cytosolic cytochrome c, cleaved caspase-3, and decreased phosphorylated Akt (Ser473), phosphorylated glycogen synthase kinase-3beta (GSK-3beta) (Ser9), and heat shock transcription factor-1. The activation of PI3K protects the cells against L-dopa neurotoxicity (Park et al. 2009). L-dopa has been reported to induce cell death in SH-SY-5Y cells with the concomitant deactivation of glutaredoxin. This compound was deactivated in a dose-dependent manner, suggesting an irreversible adduction of L-dopachrome to its nucleophilic active-site Cys-22. L-dopa also decreases thioredoxin and thioredoxin reductase activity and protein content. Knockdown of glutaredoxin increases L-dopa-induced apoptosis (Sabens et al. 2010). L-dopa activates apoptosis signaling kinase 1 (ASK1), as indicated by the phosphorylation of its downstream mitogen-activated protein kinases (MAPK), p38 and JNK. The inhibition of either p38 or JNK or the knockdown of ASK1 provides protection against L-dopa-induced apoptotic neuronal cell death in the SH-SY-5Y cell line (Liedhegner et al. 2011). L-dopa has also been found to impair proteasome activity through the  $D_1$  dopamine receptor (Berthet et al. 2012). The ability of L-dopa to be neurotoxic is dependent on its ability to oxidize to L-dopa o-quinone, which



**Fig. 4** *Possible metabolism of cyclized L-dopa o-quinone*. L-dopa oxidizes to L-dopa o-quinone, which cyclizes to cyclized L-dopa o-quinone. This compound is able to (i) polymerize to form neuromelanin, (ii) form adducts with proteins, (iii) form adducts with biomolecules, (iv) be oneelectron reduced to the cyclized L-dopa o-semiquinone radical, (v) be conjugated by GSH to 4-glutathionyl-5,6-dihydroxyindoline-2-carboxylic acid, or (vi) be two-electron reduced, catalyzed by DT-diaphorase

spontaneously cyclizes to cyclized L-dopa *o*-quinone at physiological pH (Baez et al. 1994; Takeshima et al. 2011; Kostrzewa et al. 2002).

The cyclized L-dopa *o*-quinone is able to participate in several reactions, such as (i) polymerization to melanin (for review, see Prota 1995) and (ii) the formation of adducts with proteins such as glutaredoxin (Sabens et al. 2010). A significant increase in cell death and the formation of quinoproteins were observed when CATH.a cells were incubated with L-dopa (Asanuma et al. 2012). (iii) The formation of adducts with molecules such as cysteine, nicotine, amphetamine, and quercetin has also been reported (Dehn et al. 2001; Claffey and Ruth 2001; Kubo et al. 2007; Caudle et al. 2007; Müller and Muhlack 2012). (iv) L-dopa can be one-electron reduced with flavoenzymes that transfer one electron and use NADH or NADPH. (iv) L-dopa can also be two-electron reduced by DT-diaphorase (Baez et al. 1994), and (v) it can be conjugated by glutathione transferase M2-2 (Baez et al. 1997) (Fig. 4).

#### 8 L-dopa-Induced Dyskinesia

L-dopa induces neurotoxicity in cell lines that are used as a model for in vitro studies of Parkinson's disease or primary cell cultures (Melamed et al. 1998; Walkinshaw and Waters 1995; Park et al. 2009; Sabens et al. 2010; Liedhegner et al. 2011). However, these neurotoxic effects in dopaminergic neurons are not observed in animal models of L-dopa, most likely as a consequence of the efficient conversion of L-dopa to dopamine. In fact, only one L-dopa metabolite is found in the literature (L-3-omethyldopa), supporting the idea that the major route of metabolism of L-dopa is its conversion to dopamine and that under normal conditions, L-dopa is not found, free or accumulated. Although L-dopa does not induce neurotoxicity of dopaminergic neurons, as observed in cell cultures, L-dopa induces dyskinesia in approximately 40 % patients with 4–6 years of treatment (Ahlskog & Muenter 2001). The mechanism for L-dopa-induced dyskinesias is very complex, but a role for serotonergic neurons has been proposed. The rapid oscillation of striatal dopamine during L-dopa treatment has been found to be required for the induction of dyskinesias (Meissner et al. 2006; Pavese et al. 2006). In L-dopa treatment of a Parkinson's disease patient, the dopaminergic neurons that are still functioning convert L-dopa to dopamine, which is stored in monoaminergic synaptic vesicles. Dopamine is released by the axon terminal, which is regulated by D<sub>2</sub> dopamine receptors and the dopamine transporter (DAT). However, serotonergic neurons express both AADC and VMAT-2 and release dopamine, but without the regulation of the  $D_2$  receptor and DAT (for review, see Cheshire and Williams 2012). Studies in animals with experimental Parkinsonism showed that serotonergic neurons are responsible for 80 % of dopamine release during L-dopa treatment (Tanaka et al. 1999)

## 9 Dopamine and L-dopa as Selective Neurotoxins

Dopamine is an essential neurotransmitter in dopaminergic neurons, accumulating inside monoaminergic synaptic vesicles for use as a neurotransmitter. Dopamine released into the intersynaptic space is recaptured by dopaminergic neurons with DAT, whereas cytosol dopamine can be (i) accumulated into monoaminergic synaptic vesicles mediated by VAMT-2, (ii) degraded by MAO and COMT, or (iii) autoxidized into aminochrome. However, dopamine induces neurotoxicity when undergoing oxidation to aminochrome, which can be neurotoxic by (i) forming adducts with proteins, such as alpha-synuclein, parkin, mitochondrial complexes I and III, isocitrate dehydrogenase, actin, tubulin, or the dopamine transporter or by (ii) being one-electron reduced by flavoenzymes that use NADH or NADPH as an electron donor. It was proposed that aminochrome is the endogenous neurotoxin that is responsible for the neurodegenerative process of neuromelanin-containing dopaminergic neurons (Lozano et al. 2010; Paris et al. 2010, 2011; Muñoz et al. 2012 a, b). Aminochrome is a selective neurotoxin, as it is formed inside a single dopaminergic neuron. Aminochrome can induce

neurotoxicity, resulting in a focalized neurotoxic event, which is in line with the slow neurodegeneration that occurs in neuromelanin-containing dopaminergic neurons, which takes years.

Experiments with cell cultures demonstrate the potential neurotoxic effects of L-dopa, but depending on the cellular model used for these experiments, it is possible that a part or all of the L-dopa added to the cell cultures was decarboxylated to dopamine before it induced neurotoxicity. It seems plausible that L-dopa is only a transient precursor in the synthesis of dopamine because the synthesis of dopamine is performed in a type of complex with TH and AADC, which are associated with VMAT-2, localized in monoaminergic synaptic vesicles. This suggests that L-dopa is not free in the cytosol because free L-dopa is immediately converted to dopamine and transported into the vesicles (Cartier et al. 2010). In Parkinson's therapy, L-dopa also acts as dopamine, as the decarboxylation of L-dopa and the unregulated release of dopamine from serotonergic neurons occur in both the peripheral and central nervous system, a process that seems to be involved in L-dopa-induced dyskinesias (Cheshire and Williams 2012).

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# Nature of DSP-4-Induced Neurotoxicity

# Aleksandra Bortel

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#### Abstract

N-(2-chloroethyl)-N-ethyl-2-bromobenzylamine (DSP-4) is a noradrenergic neurotoxin which selectively damages noradrenergic projections originating from the locus coeruleus (LC) and transiently alters sympathetic neurons in the periphery. DSP-4 accumulates intraneuronally and produces nerve terminal degeneration via alkylation of diverse neuronal structures. DSP-4 inhibits nor-epinephrine (NE) reuptake, stimulates NE release, and increases turnover rate of NE. Systemic administration of DSP-4 has a rapid dose-dependent depleting effect on the endogenous NE level. The effect of DSP-4 is largely restricted to

A. Bortel

Departments of Neurology, Neurosurgery, and Biomedical Engineering, Montreal Neurological Institute, McGill University, Montreal, QC, Canada e-mail: aleksandra.bortel@mcgill.ca

noradrenergic neurons; however, DSP-4 slightly reduces the level of 5-hydroxytryptamine and dopamine. The rat age at the time of DSP-4 injection is important in determining the nature of the long-term changes in the noradrenergic system. Thus, DSP-4 treatment of adult rats produces morphological changes of NE neurons with a pronounced decrease in NE levels in the cerebral cortex, hippocampus, spinal cord, and cerebellum. Less affected are the hypothalamus and pons-medulla. This process is not long lasting and after several months a regeneration of NE nerve terminals is observed. DSP-4 injected to newborn rats induces an alteration of the postnatal development of noradrenergic system with a permanent NE denervation in brain areas distal to LC cell bodies (cortex and hippocampus) and a hyperinnervation in regions proximal to LC (brainstem, cerebellum, and pons-medulla). This type of hyperinnervation does not exist after DSP-4 treatment of adult animals. The noradrenergic lesion obtained with DSP-4 is highly reproducible; therefore, DSP-4 may represent a suitable tool to discern LC neuron degeneration and recovery and to investigate the projections of non-coerulean NE neurons.

#### Keywords

Degeneration • Depletion • Dopamine- $\beta$ -hydroxylase • DSP-4 •		
Hyperinnervation • Locus coeruleus • Monoamine oxidase • Neurotoxin •		
Non-coerulean innervation • Noradrenergic neurons • Noradrenergic receptors •		
Norepinephrine transporter • Norepinephrine • Tyrosine hydroxylase		

#### List of Abbreviations

	Jeviations -
5-HIAA	5-hydroxyindoleacetic acid
5-HT	Serotonin
AD	Aldehyde dehydrogenase
AR	Aldehyde reductase
CNS	Central nervous system
COMT	Catechol-O-methyl transferase
DA	Dopamine
DAT	Dopamine transporter
DBH	Dopamine-β-hydroxylase
DHPG	3,4-dihydroxyphenylethylene glycol
DOPAC	3,4-dihydroxyphenylacetic acid
DSP-4	N-(2-chloroethyl)-N-ethyl-2-bromobenzylamine
GABA	γ-aminobutyric acid
i.p.	Intraperitoneal injection
i.v.	Intravenous injection
LC	Locus coeruleus
MAO	Monoamine oxidase
MHPG	3-methoxy-4-hydroxyphenylethylene glycol
NE	Norepinephrine
NET	Norepinephrine transporter

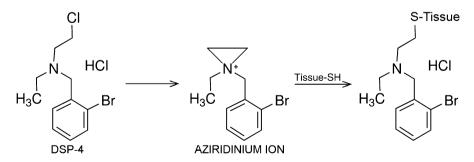
OCTs	Organic cation transporters
s.c.	Subcutaneous injection
SERT	Serotonin transporter

#### 1 Introduction

N-(2-chloroethyl)-N-ethyl-2-bromobenzylamine (DSP-4) was originally described by Ross (1976) who reported its effects on peripheral and central norepinephrine (NE) tissue levels. DSP-4 is a noradrenergic neurotoxin which crosses the blood-brain barrier and selectively damages noradrenergic projections originating from the locus coeruleus (LC). DSP-4 is a 2-chloroethylamine that interacts with the NE transporter (NET), whereby DSP-4 is then accumulated intraneuronally to induce degeneration of norad-renergic terminals (Ransom et al. 1985; Dudley et al. 1990; Howard et al. 1990). DSP-4 administration into rodents permanently inactivates NE reuptake, stimulates NE release, increases NE turnover rate in some brain regions, and produces morphological changes in NE cells (Jaim-Etcheverry 1998; Prieto and Giralt 2001).

# 2 Molecular Mechanism of Action

DSP-4 undergoes intramolecular cyclization to form an aziridinium ion, an electrophilic intermediate that can subsequently hydrolyze or react with other nucleophiles, including those present in biological tissues (thiols or amines) (Fig. 1). The aziridinium ion is a structural analogue of bretylium that is a substrate for the NET. Thus, the aziridinium ion covalently bonds with electrophilic centers on or near the site of action, disrupts the function of the NET binding site, and thereby inhibits NE reuptake (Ransom et al. 1982;



**Fig. 1** DSP-4 cyclization and reaction with nucleophiles present in biological tissues (Based on Dudley et al. 1990)

Dudley et al. 1990; Jaim-Etcheverry 1998). Inactivation of the transport system is attributable to DSP-4 alkylation at this site (Lee et al. 1982; Hallman and Jonsson 1984; Dudley et al. 1990) and requires a functional transporter to inhibit NE reuptake (Dudley et al. 1990). DSP-4 in aqueous solutions has a half-life of 7 min at pH 7.4 and 37 °C.

# 3 DSP-4 Neurotoxic Effects

#### 3.1 DSP-4 Specificity and Monoamine Reuptake

DSP-4 treatment, while having no major influence on NE cell bodies, dendrites, and preterminal axons, exerts neurotoxic effects on the terminal part of NE axons (Jonsson et al. 1981; Fritschy and Grzanna 1989, 1991a). Moreover, the part of NE axons along their pathways positioned between its origin and its entry into a terminal field is not sensitive to DSP-4. Complete loss of NE axons is found in brain regions such as the cerebral cortex, hippocampus, tectum, cerebellum, and spinal cord dorsal horn, and any changes are noted in the basal forebrain, hypothalamus, and some brainstem nuclei (Fritschy and Grzanna 1989, 1991a). Thus, DSP-4 has a preferential, selective neurotoxic effect on nerve terminal projections originating from NE perikarya of LC and does not affect NE axons in brain regions receiving a dense innervation from non-coerulean NE cells (Jonsson et al. 1981; Fritschy and Grzanna 1989, 1991a). Non-coerulean neurons provide the major NE innervation to the regions involved in the regulation of autonomic and motor activities. In contrast, LC neurons innervate brain regions involved in the processing of sensory inputs (Fritschy and Grzanna 1989, 1991a). DSP-4 initially induces degeneration of LC axon terminals that in the long term is responsible for LC cell death. The delayed death of LC neurons is probably a consequence of the loss of terminal axons rather than a result of a direct effect of DSP-4 on LC cell bodies (Fritschy and Grzanna 1991b). Initially, it was suggested that DSP-4 specificity to the noradrenergic projections originating from LC depends on the pharmacologically distinct affinity of DSP-4 for the NET in LC and non-coerulean NE axons (Fritschy and Grzanna 1989, 1991a). Subsequently, Zaczek et al. (1990) proposed that NET binding sites have distinct kinetic characteristics in the cerebral cortex (receiving LC innervation) and hypothalamus (receiving non-coerulean innervation), and therefore, DSP-4 shows a higher affinity for NET in LC versus non-coerulean nerve terminals. There are two components of NE reuptake, and only the high-affinity sites of NET are abolished by DSP-4 treatment. Low-affinity sites remain unaffected by DSP-4 (Lee et al. 1982; Hughes and Stanford 1998a). However, several studies deny NET heterogeneity and demonstrate that NETs are similar in all brain regions (Snyder and Coyle 1969; Ghraf et al. 1983). More plausibly, there may be greater intra-axonal DSP-4 accumulation in LC than noncoerulean NE axons. Interaction of DSP-4 with NET is a prerequisite for NE depletion because the NET is responsible for intracellular accumulation of DSP-4 or its aziridinium ion (Landa et al. 1984). Thus, the cytotoxic DSP-4 effect is not due to DSP-4 affinity to the transporter but depends on DSP-4 accumulation within the cells by the NET, which then disables DSP-4 outward transport. The outward transport becomes inhibited and DSP-4 remains intracellularly trapped. DSP-4 ultimately leads to ATP depletion and other cytotoxic reactions (Wenge and Bönisch 2009). NE storage is ATP dependent; therefore, DSP-4 inhibition of ATP synthesis results in NE depletion (Dudley et al. 1990).

Neuronally released monoamines are removed from the synaptic cleft by rapid reuptake into presynaptic neurons via Na<sup>+</sup>- and Cl<sup>-</sup>-dependent neuronal monoamine transporters such as NET. Additionally, monoamines are accumulated in extraneuronal tissues by Na<sup>+</sup>- and Cl<sup>-</sup>-independent extraneuronal monoamine transporters and organic cation transporters (OCTs). These transporters are involved in the absorption, distribution, and elimination of endogenous compounds (Wenge and Bönisch 2009). DSP-4 interacts at the level of neuronal membrane and forms covalent bonds with thiol groups of proteins (Jaim-Etcheverry and Zieher 1980; Dudley et al. 1990). DSP-4 and aziridinium ion are competitive inhibitors of NE reuptake, while due to the alkylation of NET, DSP-4 blocks the transporter irreversibly (Zaczek et al. 1990). DSP-4 also inhibits other transporters such as that for dopamine (DAT) and serotonin (SERT). About 80 % and 60 % of the transport activity of DAT and SERT, respectively, is inactivated by the neurotoxin. DSP-4 inhibits, as well, OCTs with a potency 3.2-fold (OCT2), 7.5-fold (OCT1), or 9-fold (OCT3) higher than for the NET. The neurotoxic DSP-4 effect is completely irreversible at NET; only partially irreversible at DAT, SERT, and OCT3; and fully reversible at OCT1 and OCT2 (Wenge and Bönisch 2009).

# 3.2 Synaptic Norepinephrine Levels and Noradrenergic Receptors

DSP-4-induced imbalance between NE release and reuptake leads to an increase in the extracellular NE concentration and NE efflux in the frontal cortex of freely moving rats (Hughes and Stanford 1998a, b). However, DSP-4 treatment does not alter the basal extracellular NE level in the frontal cortex of anesthetized rats, although NE release evoked by administration of an  $\alpha_2$ -adrenergic receptor antagonist results in lower but still significant increase in extracellular NE levels (Kask et al. 1997; Harro et al. 1999). The same, lower but still significant increase in NE release induced by a GABA<sub>A</sub> antagonist (bicuculline) was observed by Giorgi et al. (2003) in the piriform cortex. Long-term NE depletion and the resulting decreased presynaptic NE release affect the function of pre- and postsynaptic NE receptors that are exposed to the environment of reduced synaptic NE concentration (Dudley et al. 1990). DSP-4 is a blocker of  $\alpha$ -adrenergic receptors; therefore, DSP-4 initially reduces a-adrenergic receptor binding. This blocking effect is not long lasting because already 14 days after the neurotoxin application  $\alpha$ - and  $\beta$ -adrenergic receptor binding recovers (Jonsson et al. 1981). DSP-4 exerts a consistent upregulation of β-adrenergic receptors in the somatosensory cortex and hippocampus at 1–3 weeks after treatment (Zahniser et al. 1986). DSP-4 likewise increases

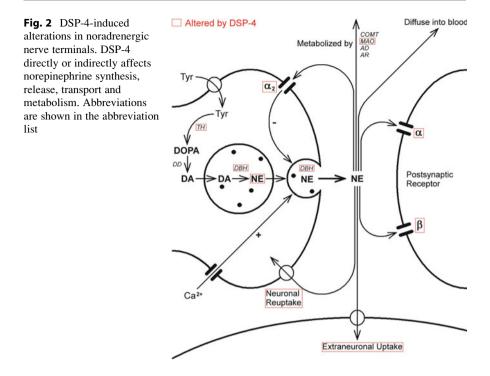
the number of  $\beta$ -adrenergic receptors and does not change the affinity of these receptors. The density of binding sites of  $\beta$ -adrenergic receptors is enhanced at 1 day, 7 days, and up to 1 year after central NE denervation induced by DSP-4 (Theron et al. 1993; Wolfman et al. 1994). The number and the affinity of  $\alpha_1$ -adrenergic receptors are unchanged in somatosensory cortex and hippocampus 1–3 weeks after DSP-4 treatment (Zahniser et al. 1986; Szot et al. 2010). As well, the properties of  $\alpha_2$ -adrenergic receptors are unaltered (Zahniser et al. 1986). Another study shows that the activity of  $\alpha_2$ -adrenergic receptors is reduced in the parietal cortex and hippocampus 3 weeks after DSP-4 treatment, but not in the hypothalamus (Prieto and Giralt 2001). Interestingly, it was reported that the number of  $\alpha_2$ -adrenergic receptors is reduced in the cortex (20 %), hippocampus (18 %), cerebellum (24 %), and hypothalamus (39 %) at 3 days after DSP-4 (100 mg/kg; DSP-4 dose is mentioned only when is different from 50 mg/kg) treatment and unchanged or even increased 15 days after treatment. Thus, decreased density of cortical presynaptic  $\alpha_2$ -adrenergic receptors is progressively masked by increased sensitivity of postsynaptic  $\alpha_2$ -adrenergic receptors (Heal et al. 1993). The density of binding sites of  $\alpha_1$ - and  $\alpha_2$ -adrenergic receptors is increased in cerebral cortex and cerebellum 7 days after DSP-4 administration, while  $\alpha_1$ - and  $\alpha_2$ -adrenergic receptor number attains control values 3 months and 1 year later, respectively (Theron et al. 1993; Wolfman et al. 1994).

The enhancement of  $\alpha$ -adrenergic receptor density is an adaptative change of postsynaptic receptors as a consequence of the long-term disruption in NE transmission. However, an irreversible increase of  $\beta$ -adrenergic receptors may be linked with the adaptative changes that are independent of the endogenous NE concentration (Dooley et al. 1987).

# 3.3 Norepinephrine Synthesizing and Metabolizing Enzymes

DSP-4 induces a profound loss of both NE and dopamine-β-hydroxylase (DBH) from noradrenergic axons, with an uncorrelated time course (Fig. 2) (Ross 1976; Jaim-Etcheverry and Zieher 1980; Fritschy et al. 1990). Tissue NE levels decline within hours, and DBH activity is reduced within 4 days after DSP-4 treatment (Fritschy et al. 1990). Therefore, two phases in the response of NE axons to DSP-4 are suggested: (1) an acute phase that is characterized by a marked loss of neuro-transmitter and (2) a neurodegenerative phase associated with loss of DBH and structural disintegration of NE axons (Fritschy et al. 1990). LC cells which survive the injury induced by DSP-4 exhibit greater DBH immunocytochemical reaction (Fritschy and Grzanna 1991b).

There are contradictory studies concerning the DSP-4 influence on tyrosine hydroxylase, a synthesizing enzyme that is localized in noradrenergic neurons and regulated independently (Fig. 2). It is proposed that DSP-4 treatment induces a decrease of basal tyrosine hydroxylase activity in the hippocampus and parietal cortex, whose axon terminals arise exclusively from LC, but not in the hypothalamus which is innervated by noradrenergic terminals originating from the tegmental lateral



nuclei (Prieto and Giralt 2001). However, other studies show that DSP-4 exerts a long-lasting NE depletion with compensatory increased accumulation of tyrosine hydroxylase in noradrenergic neurons (Booze et al. 1988; Dudley et al. 1990).

It was suggested that DSP-4 may also interfere with some enzymes involved in NE metabolism. First, DSP-4 inhibits monoamine oxidase (MAO) activity in peripheral tissues (Lyles and Callingham 1981). Furthermore, the MAO-B inhibitor deprenyl effectively blocks the neurotoxic action of DSP-4 on noradrenergic neurons. However, MAO-B inhibition of DSP-4 toxicity is unrelated to NE metabolism, because MAO-A plays a central role in metabolizing endogenous monoamines including NE (Gibson 1987). Finnegan et al. (1990) have proposed that deprenyl neuroprotective effects are unrelated to MAO-B inhibition but to other properties such as the effects on the monoamine release, reuptake and metabolism, as well as the effects on other central neurotransmitter systems. Indeed, DSP-4 or aziridinium ion binds to the active catalytic site of MAO-B but is not further metabolized. Therefore, the neuroprotective deprenyl action depends on the blockade of DSP-4 reuptake into noradrenergic neurons (Dudley et al. 1990; Finnegan et al. 1990). However, the neuroprotective action of highly potent and selective MAO-B inhibitors, that do not interfere with DSP-4 reuptake within NE terminals (as it is observed with deprenyl), likewise fully prevents DSP-4-induced NE depletion (Yu et al. 1994; Zhang et al. 1996). Conversely, some selective MAO-B inhibitors are inactive against DSP-4-induced neurotoxicity (Bertocci et al. 1988; Finnegan et al. 1990), and the absence of MAO-B (i.e., MAO-B knockout mice) does not influence the mechanism of neurotoxic DSP-4 action (Fornai et al. 2001). Therefore, the neuroprotective and/or neurorescue effects of some MAO-B inhibitors are unclear.

### 3.4 Electrophysiological Responsiveness

DSP-4 increases NE turnover in brain regions in which NE depletion is greater than 85 % (cerebral cortex and hippocampus), but has no effect on NE turnover when NE is depleted by only 30–60 % (cerebellum, hypothalamus, brainstem, and LC). Firing rates of monoaminergic neurons increase only after 75 % NE depletion. Thus, increased firing represents a compensatory response to the injury, resulting in increased transmitter release from the terminals spared by the lesion (Chiodo et al. 1983; Logue et al. 1985). Olpe et al. (1983) reported that DSP-4 decreases the mean neuronal LC firing rate without altering the structural appearance of LC perikarya. Precisely, DSP-4 does not affect the basal activity of LC noradrenergic neurons but increases the irregularity and the burst firing of noradrenergic neurons 2 weeks after DSP-4 treatment (Szot et al. 2010).

# 4 DSP-4 Doses and Routes of Administration

Systemic administration (s.c., i.p. or i.v.) of DSP-4 has a rapid dose-related effect on tissue levels of NE (Table 1). Effects are achieved with a DSP-4 dose of 20–100 mg/kg i.p. by 0.5 h after treatment (Hallman and Jonsson 1984; Jaim-Etcheverry 1998). Newborn rats are typically treated s.c., while adult rats are routinely given i.p or i.v. injections. Notably, NE depletion is much greater after i.v. than after i.p. injection of adult rats (Jaim-Etcheverry and Zieher 1980); however, i.p. administration is sufficient to produce pronounced NE depletion (up to 95 %). A dose of 50 mg/kg of DSP-4 is the most efficient in producing pronounced, long-lasting reduction of NE reuptake in the CNS without causing any lethal effects in rats (Jonsson et al. 1981) and mice (Dailly et al. 2006).

Neonatal DSP-4 treatment leads to dose-dependent adulthood changes of NE concentrations in assorted brain regions. The lowest tested DSP-4 dose (10 mg/kg s.c.) produces about 50 % NE reduction in cerebral cortex and spinal cord and small changes in cerebellum and pons-medulla. A large DSP-4 dose (50 mg/kg s.c.) produces near complete NE depletion in cerebral cortex and spinal cord and associated increases in NE concentration, both in the cerebellum and pons-medulla (Jonsson et al. 1982). The latter effect is reflective of reactive collateral sprouting of bi- or multipolar NE neurons of LC, following DSP-4 injury of NE terminals and/or axons projecting to cerebral cortex and hippocampus – as was found in similar studies with the neurotoxins 6-hydroxydopamine (Sachs and Jonsson 1975) and 6-hydroxydopa (Kostrzewa and Garey 1977).

DSP-4 doses	<25 mg/kg	25 mg/kg	50 mg/kg	100 mg/kg
Cortex	-	_	X	X
NE degeneration				
Spinal cord	_	Х	Х	X
NE degeneration				
Brainstem	_	Х	Х	_
NE degeneration (A)				
NE hyperinnervation (N)				
Cerebellum	Х	Х	Х	_
NE degeneration (A)				
NE hyperinnervation (N)				

**Table 1** Dose-dependent changes of NE tissue levels in different brain areas observed 6 weeks after DSP-4 treatment (Based on Jaim-Etcheverry and Zieher 1980)

(A) NE degeneration observed after DSP-4 injection to adult animals, and (N) NE hyperinnervation after DSP-4 injection to newborn animals

# 5 The Nature and Extension of DSP-4-Induced Neurotoxicity in Adulthood

DSP-4 treatment does not have significant sex differences with respect to the regional alterations in the monoamine levels (Jonsson et al. 1981, 1982). Also, body weight gain in both sexes is virtually identical after DSP-4 and similar to that of control litters. As adults, the weight of neonatally treated rats of both sexes is approximately 95 % of the weight of age-matched controls (Jonsson et al. 1981, 1982). The age of rats at the time of DSP-4 treatment is critical in determining the nature and extent of the long-term changes in noradrenergic neurons (Jaim-Etcheverry 1998).

# 5.1 DSP-4 Treatment of Adult Rats

#### 5.1.1 Noradrenergic System

DSP-4 treatment of adult rats generally causes a decrease of tissue levels of NE and its two principal metabolites: 3,4-dihydroxyphenylethylene glycol (DHPG) and 3-methoxy-4-hydroxyphenylethylene glycol (MHPG) (Dudley et al. 1990; Hallman et al. 1984; Logue et al. 1985). However, this process is not long lasting, and after several months a regeneration of NE nerve terminals is observed (Hallman et al. 1984). DSP-4 neurotoxic effects are rapid, being maximal 2 h after DSP-4 injection for NE reuptake in vitro and by 4–6 h for endogenous NE levels. Maximal NE depletion is achieved most rapidly in the occipital cortex, hippocampus, hypothalamus, pons-medulla, cerebellum, and spinal cord, but later, in the frontal cortex, striatum, and mesencephalon. Acutely, DSP-4 has a profound NE-depleting effect in the cortex and cerebellum, but only a modest effect in ventral forebrain and hypothalamus (Grzanna et al. 1989). Simultaneously, a profound reduction in NE axon staining is found in the cerebral cortex and cerebellum after DSP-4 treatment, whereas the ventral forebrain and hypothalamus appear to be unaffected (Grzanna et al. 1989). Ten days after DSP-4 treatment, NE concentrations are decreased by about 90 % in the cerebral cortex, hippocampus, spinal cord, and cerebellum. Less affected are the hypothalamus and pons-medulla, known to have a mixed NE innervation (both from LC and other noradrenergic nuclei in brainstem). DSP-4 decreases endogenous tissue levels of NE, not only at noradrenergic nerve terminals but also at both noradrenergic and serotoninergic cell body levels. Seven days after DSP-4, NE concentrations are decreased in LC perikarya and dorsal raphe nucleus (Cassano et al. 2009). Thus, DSP-4 neurotoxicity is exerted on the LC via retrograde degeneration (Fritschy and Grzanna 1991a) as well as from direct DSP-4 action at the LC itself, where NET is highly expressed (Cassano et al. 2009). Alterations in NE reuptake in assorted brain areas are compatible with observed changes in tissue levels of NE after DSP-4 (Jonsson et al. 1981).

NE depletion induces profound adaptive changes in other neuronal phenotypes. DSP-4 does not modify endogenous levels of brain acetylcholine,  $\gamma$ -aminobutyric acid (GABA), glutamic acid, glycine, or aspartic acid. However, DSP-4 depletes tissue levels of 5-HT and slightly depletes DA (Jaim-Etcheverry and Zieher 1980; Jonsson et al. 1981; Fornai et al. 1996b).

#### 5.1.2 Serotoninergic System

DSP-4 induces an acute decrease of 5-HT concentrations in the cerebral cortex, mesencephalon, and spinal cord (Jackisch et al. 2008), and at 8 weeks 5-HT is depleted as much as 85 % in the cerebral cortex (Jonsson et al. 1981). DSP-4 does not alter the level of 5-HT and its metabolite 5-hydroxyindoleacetic acid (5-HIAA) in hippocampus (Jackisch et al. 2008) and dorsal raphe nucleus (Cassano et al. 2009). However, one day after DSP-4 treatment, a prominent increase in 5-HIAA is observed in the cortex, hippocampus, and hypothalamus (Theron et al. 1993). Because NE normally inhibits presynaptic 5-HT release (Vizi and Kiss 1998), noradrenergic denervation of hippocampus is associated with an increase in basal and evoked 5-HT release and reduced 5-HT accumulation in the hippocampal slices (Cassano et al. 2009). The neurotoxic effect of DSP-4 on 5-HT neurons can be prevented by pretreatment with the SERT blocker (zimelidine) (Jonsson et al. 1981).

#### 5.1.3 Dopaminergic System

DSP-4 alters DA levels in several brain regions (Jonsson et al. 1982). Thus, DSP-4-induced noradrenergic lesioning reduces DA levels in hippocampus, cerebellum, prefrontal cortex, LC, and dorsal raphe nucleus (Jonsson et al. 1981; Cassano et al. 2009). Moreover, DSP-4 treatment decreases (Lategan et al. 1992) or does not change (Hatip-al-Khatib et al. 2001; Häidkind et al. 2002) the extracellular DA concentration and reduces the extracellular 3,4-dihydroxyphenylacetic acid (DOPAC) level in nucleus accumbens (Lategan et al. 1992; Hatip-al-Khatib et al. 2001). A reduction in extracellular DA concentration is attributable to a reduction in basal DA release (Lategan et al. 1992; Weinshenker et al. 2008).

#### 5.1.4 Other Neuronal Phenotypic Systems

The hippocampal endogenous acetylcholine level (Hörtnagl et al. 1989) and activity of the acetylcholine transporter (Harrell et al. 2005) are unaltered following DSP-4 treatment. However, DSP-4-induced lesion of noradrenergic neurons facilitates cholinergic transmission in the hippocampus (Jackisch et al. 2008). DSP-4 treatment likewise induces an upregulation of the turnover and activity of cholinergic neurons. Acetylcholine accumulation is enhanced in the hippocampus, while basal and evoked acetylcholine efflux are increased in DSP-4-treated rats (Jackisch et al. 2008).

#### 5.1.5 Peripheral Sympathetic Neurons

Related DSP-4 effects are observed in peripheral sympathetic neurons as compared with central NE neurons. However, a complete recovery of peripheral adrenergic nerves is observed 1 week after DSP-4, and a gradual recovery of decreased NE tissue levels and reuptake occurs by 2–4 weeks after DPS-4 treatment (Jaim-Etcheverry et al. 1980; Jonsson et al. 1981).

### 5.2 DSP-4 Treatment of Newborn Rats

#### 5.2.1 Noradrenergic System

Systemic DSP-4 treatment of newborn rats results in an alteration in postnatal development of the noradrenergic system, with a permanent denervation of distal NE projection regions and associated NE hyperinnervation in regions proximal to LC cell bodies. The total number of NE nerve terminals is considered to be unaltered (Jaim-Etcheverry and Zieher 1980). NE hyperinnervation is considered to be a consequence of collateral NE accumulation and/or sprouting of noradrenergic neurons (Jaim-Etcheverry and Zieher 1980). Neonatal DSP-4 treatment leads to adulthood changes of tissue NE levels in several brain regions. In adulthood NE is depleted approximately 90 % in cerebral cortex, olfactory bulb, hippocampus, and spinal cord of rats treated as neonates with DSP-4 (Jonsson et al. 1982; Nowak et al. 2006; Bortel et al. 2008b), with less pronounced NE effects in hypothalamus and septum. No significant effects are noted in the striatum, olfactory tubercle, and medulla oblongata (Jonsson et al. 1982; Nowak et al. 2006), while a pronounced increase in NE concentrations are present in the cerebellum, brainstem, mesencephalon, and pons-medulla after DSP-4 (Jonsson et al. 1982; Bortel et al. 2008b). It is notable that NE concentrations are decreased in the cerebellum, mesencephalon, and pons-medulla 6 h after DSP-4 administration, in contrast to the marked increase in NE tissue levels in these regions in adulthood (Jonsson et al. 1982).

When administered at birth, DSP-4 decreases in vitro NE reuptake in cerebral cortex and spinal cord by approximately 90 %, but does not alter NE reuptake in the striatum and increases NE reuptake in the cerebellum (Jonsson et al. 1982).

Permanent NE denervation in cerebral cortex and spinal cord is present at all developmental stages, although more pronounced when animals are treated with DSP-4 up to the 5th day after birth. However, NE hyperinnervation in the cerebellum

and pons-medulla is only observed when DSP-4 is administered to 1 and 3 days old rats (Jonsson et al. 1982; Bortel et al. 2008b). By the age of 8 days, permanent NE depletion is present in these structures of the brain of animals treated at birth with DSP-4 (Jonsson et al. 1982). Acutely (1 day after DSP-4 treatment) NE depletion is observed in animals of all ages (Jonsson et al. 1982).

#### 5.2.2 Serotoninergic System

DSP-4 induces dose-dependent changes of 5-HT tissue levels in several brain regions. DSP-4 reduces the endogenous 5-HT concentrations in the cerebral cortex and spinal cord and reduces 5-HT reuptake in the occipital cortex, hippocampus, and cerebellum. No significant effects are noted in other brain regions (Jonsson et al. 1981).

Neonatal DSP-4 treatment does not induce any change in 5-HT levels and 5-HT reuptake when whole brain is analyzed (Jonsson et al. 1982). SERT blocker zimelidine attenuates the effects of DSP-4 on 5-HT and 5-HIAA in the frontal cortex, hippocampus, striatum, and hypothalamus (Jonsson et al. 1981; Dabrowska et al. 2007).

#### 5.2.3 Dopaminergic System

Although neonatal DSP-4 treatment does not acutely (i.e., directly) affect DA and DOPAC levels in the hippocampus and striatum (Nowak et al. 2006), there is an adaptive decrease in DA content of occipital cortex, hippocampus, cerebellum, mesencephalon (Jonsson et al. 1982), and dorsal raphe nuclei (Cassano et al. 2009) in adulthood.

#### 5.2.4 GABAergic System

Neonatal DSP-4 treatment of rats does not alter GABA tissue levels in prefrontal cortex, hippocampus, brainstem, and cerebellum (Bortel et al. 2008b) and does not alter the extracellular GABA concentration in prefrontal cortex (Bortel et al. 2008a). However, DSP-4-induced noradrenergic denervation of the cerebral cortex reduces the number of GABA<sub>A</sub> receptors that are localized on the presynaptic axons and nerve terminals of noradrenergic neurons (Medina and Novas 1983; Suzdak and Gianutsos 1985). This significant decrease in density of benzodiazepine receptors persists even 1 year after DSP-4 treatment (Wolfman et al. 1994). Moreover, DSP-4 increases the number of benzodiazepine receptors in the cerebellum and brainstem (Medina and Novas 1983).

#### 5.2.5 Peripheral Sympathetic Neurons

The effect of DSP-4 on the peripheral sympathetic nervous system is minor. DSP-4 fails to permanently alter the NE concentration in peripheral organs (Jaim-Etcheverry and Zieher 1980; Jonsson et al. 1982). At 1 month after DSP-4 treatment, NE tissue levels approximate control values (Jonsson et al. 1982). Neonatal DSP-4 treatment does not produce long-lasting changes in NE content of heart, salivary glands, and spleen (Jaim-Etcheverry and Zieher 1980; Jonsson et al. 1981).

#### 5.3 Prenatal DSP-4 Treatment

Prenatal DSP-4 administration (at the end of gestation; twice 25 mk/kg i.v.) modifies NE tissue levels to the same extent as when DSP-4 is administered to newborn animals (Jaim-Etcheverry and Zieher 1980). DSP-4 prenatal treatment permanently decreases NE levels in cerebral cortex and spinal cord, demonstrating that the neurotoxin passes the blood-placenta barrier and has a potent neurotoxic effect in the prenatal stage. Interestingly, there is no ensuing increase in NE concentration in the cerebellum and pons-medulla of treated prenates when observed as adults – in contrast to effects after DSP-4 treatment of newborn animals (Jonsson et al. 1981).

# 6 Rodent Differences in the Susceptibility to DSP-4

DSP-4-induced neurodegenerative changes in noradrenergic neurons in the CNS are qualitatively similar in both rats and mice (Jonsson et al. 1981). However, NE depletion in forebrain regions such as cortex and hippocampus appears to be more pronounced and persistent in mice versus rats (Fornai et al. 1996a; Dailly et al. 2006; Cassano et al. 2009; Szot et al. 2010). As well, sympathetic adrenergic nerves of the mouse are more sensitive than rats to the neurotoxic action of DSP-4. This effect is likely due to the differences in diffusion conditions or size of NE varicosities (Jonsson 1980). It is notable that various mouse and rat strains show a different sensitivity in the CNS to the neurotoxic DSP-4 action, as exemplified by the finding that Long-Evans rats are less sensitive than Sprague–Dawley rats to DSP-4 (Schuerger and Balaban 1995).

# 7 Conclusion

DSP-4 is an alkylating agent with high specificity for NE and 5-HT neurons in vivo. The neurotoxin transiently alters sympathetic neurons in the periphery and produces persistent alterations in the central noradrenergic system. DSP-4, having high affinity for the NET, and being accumulated within NE nerve endings, depletes monoamines by reducing mitochondrial function and ultimately destroying noradrenergic neurons in the periphery and brain (Jaim-Etcheverry and Zieher 1980; Dudley et al. 1990). Moreover, DSP-4 is able to cross the blood-placenta barrier in pregnant animals, thereby enabling DSP-4 to alter noradrenergic neuronal development in early stages of ontogeny.

DSP-4 has the capacity to produce destruction of NE axons in brain regions innervated by LC, without affecting non-coerulean NE axons (Fritschy and Grzanna 1989). Therefore, DSP-4 appears to be a useful denervation tool to study the projections of non-coerulean NE neurons in isolation. Additionally, DSP-4 may serve as a useful experimental tool to discern LC neuron degeneration and recovery. The long-term NE depletion and irreversible inhibition of neuronal reuptake induced by DSP-4 can be used in pharmacological investigations. Thus, the alterations in NE tissue levels and release may be used to investigate synaptic biochemistry including mechanisms of noradrenergic receptor up- or downregulation. The NE-depleting DSP-4 effects can be used to study behavioral disorders. Until now, we know that DSP-4 treatment decreases exploratory activity in rats that results from increased anxiety (Delini-Stula et al. 1984; Harro et al. 1995). DSP-4-induced NE depletion exerts deficits in working memory without affecting reference memory (DSP-4 did not affect the previously learned pattern) (Ohno et al. 1993; Sontag et al. 2008). Moreover, NE possesses an anticonvulsant effect; therefore, central NE depletion facilitates induction of seizures evoked by bicuculline, pentylenetetrazol, or maximal electroshock (Mishra et al. 1994; Giorgi et al. 2003, 2006).

The noradrenergic lesion obtained with DSP-4 is remarkably reproducible and the usefulness and impact of DSP-4-lesioned models have enormous pharmacological potential. Despite these obvious successes, limitation still exists for DSP-4 because of unspecificity in its mechanism of action. DSP-4 interacts not only with NE reuptake systems, but alters, as well, other monoamine reuptake systems as well as enzymes involved in NE synthesis. Adaptive changes in noradrenergic receptors, relative to NE depletion, further complicate the actions of DSP-4. Therefore, all experiments based on DSP-4 should be carefully interpreted.

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# MPTP Neurotoxicity: Actions, Mechanisms, and Animal Modeling of Parkinson's Disease

# Livia Pasquali, Elena Caldarazzo-lenco, and Francesco Fornai

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#### Abstract

The study of neurotoxicity induced by MPTP led to drastically change the perspective on Parkinson's Disease. In fact the selective neurotoxicity induced by MPTP rejuvenated PD research and generated a number of studies aimed at

L. Pasquali • E. Caldarazzo-Ienco

Clinical Neurology, University of Pisa, Pisa, Italy e-mail: lpasquali@med.unipi.it; elenacaldarazzoienco@gmail.com

F. Fornai (🖂)

Neuromed, I.R.C.C.S, Pozzilli, IS, Italy e-mail: francesco.fornai@med.unipi.it

Human Anatomy, University of Pisa, Pisa, Italy

elucidating the mechanisms of action of MPTP. Remarkably, these molecular mechanisms turned out to be critical also for the survival of DA neurons in idiopathic PD. In this chapter we report the main concepts developed over the last three decades to understand key molecular steps which are pivotal in MPTP toxicity. This is the case of the role played by DAT and VMAT-2 in conditioning the sensitivity to MPTP neurotoxicity. Similarly, the mitochondria as targets of MPTP toxicity appear similarly affected by selective mutation of genes leading to PD. Again, the fate of mitochondria and the ability to clear these organelles when being dysfunctional is key in the modulation of MPTP toxicity. This also applies for misfolded proteins such as alpha synuclein. Again, multiple brain areas as well as peripheral sites are increasingly recognized to be affected both during MPTP toxicity and sporadic PD patients. Nowadays it seems that MPTP per se did not lead to discovery of the environmental compound which causes PD, nonetheless the study of MPTP did disclose several molecular and cellular pathways which are critical in the genesis of PD. This latter point fairly corresponds to what we enthusiastically expected from MPTP when it was identified as a causal agent of what it remains, a toxic form of environmental parkinsonism.

#### **Keywords**

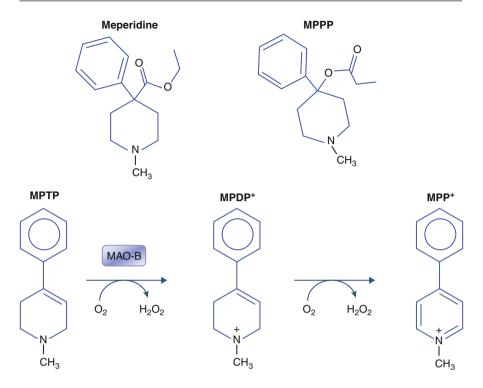
Animal models • MPP+ • Parkinsonism

List of Abbreviations		
AMPA	Alpha-amino-3-hydroxy-5-methylisoxazole-4-propionic acid	
ATG	Autophagy	
ATP	Adenosine triphosphate	
BBB	Blood–brain barrier	
COX	Cyclooxygenase	
DA	Dopamine	
DAT	DA transporter	
DSP-4	N-(-2-chloroethyl)-N-ethyl-2-bromobenzylamine	
EAA	Excitatory amino acids	
ENS	Enteric nervous system	
GI	Gastrointestinal	
GSH	Glutathione	
i.c.v.	Intracerebroventricular	
IP <sub>3</sub>	Inositol(1,4,5)trisphosphate	
JNK	c-Jun N-terminal kinase	
КО	Knockout	
LC	Locus coeruleus	
MAO-B	Monoamine oxidase type B	
METH	Methamphetamine	
MK-801	Dizocilpine	
Mn-SOD	Manganese superoxide dismutase	
$MPDP^+$	1-Methyl-4-phenyl-2,3-dihydropyridine	
$MPP^+$	1-Methyl-4-phenylpyridinium	

MPPP	1-Methyl-4-phenyl-4-propionoxy-piperidine
MPT	Mitochondrial permeability transition
MPTP	1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine
NAD	Nicotinamide adenine dinucleotide
NE	Norepinephrine
NET	NE transporter
NMDA	<i>N</i> -methyl-D-aspartate
NO	Nitric oxide
PD	Parkinson's diseases
ROS	Reactive oxygen species
SC	Spinal cord
SNpc	Substantia nigra pars compacta
SOD	Superoxide dismutase
TH	Tyrosine hydroxylase
UP	Ubiquitin-proteasome
VMAT-2	Vesicular monoamine transporter type 2

# 1 Introduction

The history of MPTP began in 1947, when researchers from a North American pharmaceutical company named Hoffmann-La Roche while developing compounds as painkillers synthesized a novel opioid derivative named meperidine (Demerol<sup>(R)</sup>). In this context, Dr. Ziering synthesized a meperidine derivative, 1-methyl-4-phenyl-4-propionoxy-piperidine (MPPP, Fig. 1), which possesses the meperidine structure, containing an esteric group (Ziering et al. 1947). Since MPPP turned out not to be more effective than Demerol, it was never marketed. In 1976, a 23-year-old Maryland chemistry graduate named Barry Kidston, in the attempt to synthesize MPPP to explore its potential as a recreational drug, esterified the intermediate tertiary alcohol of MPPP with propionic anhydride at a drastically high temperature. This led to the synthesis of 3 % of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) as a byproduct of MPPP (Fig. 1). Within 3 days following MPTP intake, Kidston developed parkinsonism, which was successfully treated with L-DOPA. However, he died 18 months later due to cocaine overdosing. The neuropathological exam was in line with neurological symptoms showing dopamine (DA) neuron loss in the substantia nigra pars compacta (SNpc) (Davis et al. 1979). Kidston's case remained unique for a few years; in 1982, in Northern California during a very short time, a sort of epidemic parkinsonism was reported in Santa Clara Valley Medical Center in San Jose (CA) by Dr. J. William Langston, the hospital's chief neurologist. Dr. Langston initially described seven cases, but this number increased soon after, when several cases of severe parkinsonism occurring in young adults were described in the same area. It was evident that these patients were drug addicts and all of them injected *intravenously* an illicit street drug called "Super Demerol"



**Fig. 1** Structural formula of meperidine analogues. The cartoon shows the structural affinity between MPTP and its metabolites with meperidine and MPPP. MPTP was produced as by-product in the effort to synthesize MPPP. Occurrence of MPTP and administration of the compound to a variety of animal species lead to the enzymatic oxidation by monoamine oxidase (MAO) type B to produce the intermediate MPDP<sup>+</sup> which, in turn, undergoes spontaneous oxidation to MPP<sup>+</sup>

which turned out to contain significant amount of MPTP (Langston et al. 1983). Also these patients showed the clinical features of idiopathic Parkinson's disease (PD), including bradykinesia, rigidity, and, in some cases, resting tremor (Ballard et al. 1985). Again, these symptoms were relieved by the intake of oral L-DOPA, while postmortem analysis confirmed the occurrence of massive cell loss in the SNpc (Ballard et al. 1985). Only later on the occurrence of Lewy bodies in these patients was demonstrated thus adding on to the remarkable overlapping with idiopathic PD (Langston et al. 1999).

The complete story of these patients was reported by Dr. William Langston in his monograph "The Case of the Frozen Addicts" (Langston and Palfreman 1996).

After identification of MPTP as the causal agent for the epidemic environmental parkinsonism, neuroscientists involved in PD research evaluated its neurotoxic effects in a variety of animal species including nonhuman primates, rodents, cats, pigs, dogs, sheep, rats, and goldfishes (Kopin and Markey 1988; Gerlach and Riederer 1996; Przedborski et al. 2001). It was soon evident that the neurotoxic properties of MPTP vary depending to which animal it was administered. In fact,

when MPTP is administered to monkeys, most clinical and pathological hallmarks of PD can be reproduced (Langston et al. 1984a; Chiueh et al. 1984; Doudet et al. 1985; Crossman et al. 1985; Nomoto et al. 1985; Vezoli et al. 2011), while rats are quite resistant to MPTP toxicity (Chiueh et al. 1984).

The discovery of selective neurotoxicity induced by MPTP rejuvenated PD research and generated a number of studies aimed at elucidating the mechanisms of action of MPTP.

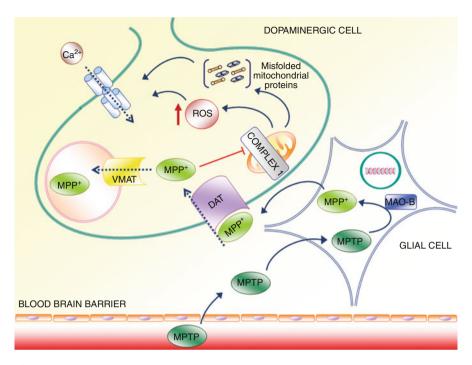
In fact, it was postulated that MPTP analogues such as pesticides may represent environmental neurotoxins responsible for the etiology of idiopathic PD. Moreover, for the first time, an experimental model mimicking both neurological symptoms and key neuropathological features of PD was available. This experimental setting appeared to be ideal to dissect the most intimate molecular mechanisms generating selective neurodegeneration underlying PD. A bulk of studies now demonstrate that MPTP analogues are not primarily involved in PD etiology. Moreover, now we are aware that PD possesses a strong genetic background which limits the role of environmental factors. Again, even in primates, some discrepancies between PD and MPTP-induced parkinsonism do exist. For instance, the striatal pattern of DA loss in PD involves more the putamen than caudate nucleus (as witnessed by pathobiochemistry, neuropathology, and more recently brain imaging, Kish et al. 1988; Jellinger 2002; Brooks and Pavese 2011), while MPTP does not discriminate between DA innervations of various neostriatal regions (Pifl et al. 1988). Nonetheless, MPTP intoxication still represents the best experimental tool to mimic very closely the selective and specific features which characterize PD. This extends also to those neuropathological features which were not immediately evident in MPTPintoxicated patients, such as the occurrence of neuronal inclusions. In fact, neuronal inclusions known as Lewy bodies which are considered a hallmark of PD were not described at first following sporadic injections of MPTP. Now we are aware that in MPTP-intoxicated patients, Lewy bodies do occur (Langston et al. 1999).

Again, clinical and pathological research on PD in the last three decades demonstrated that despite the multiple etiology (various genetic forms or idiopathic), the disease is constantly characterized by degeneration of multiple areas both within the CNS (Braak et al. 2003) and outside the brain, the spinal cord, and periphery. Accordingly, research on MPTP intoxication cops quite perfectly with such novel findings in PD. In fact, nowadays we are aware that by selecting the appropriate animal species and MPTP dosing regimen, we can reproduce most of these extra-nigral degenerations in multiple brain and peripheral areas. Thus, apart from discrepancies in the topography of nigral cell loss and nigrostriatal denervation, at present we can firmly establish the occurrence of neuronal inclusions and the involvement of multiple brain sites and peripheral regions following MPTP intoxication. This renders the MPTP model constantly updated in following progress in PD research. In fact, some potential discrepancies were resolved progressively adapting in-depth studies on MPTP intoxication to the extended degeneration which was increasingly described in PD.

In order to analyze MPTP toxicity, it is mandatory to dissect those molecular steps which characterize MPTP metabolism and the dynamics of MPTP in target cells. At the same time, after reviewing main concepts which characterize MPTP toxicity at neuronal level, we will mention briefly those neurotransmitter and biochemical pathways which either contribute to detrimental effects or mediate compensatory mechanisms which counteract MPTP intoxication.

# 2 MPTP Metabolism

MPTP is a highly lipophilic substance (Fig. 1) that easily crosses the blood-brain barrier (BBB) (Fig. 2). In the brain, MPTP is metabolized to 1-methyl-4phenylpyridinium (MPP<sup>+</sup>) (Figs. 1 and 2) via the formation of the intermediate 1-methyl-4-phenyl-2,3-dihydropyridine (MPDP<sup>+</sup>, Fig. 1), this oxidation being catalyzed by monoamine oxidase type B (MAO-B) (Chiba et al. 1984: Kopin 1987) (Figs. 1 and 2). Once formed, MPP<sup>+</sup>, which is a polar compound, cannot cross back the BBB. MPP<sup>+</sup> is the real neurotoxin at cellular level; in fact MAO inhibitors (which impede the generation of MPP<sup>+</sup> from MPTP) provide complete protection against MPTP toxicity (Heikkila et al. 1984a; Langston et al. 1984b). When MPP<sup>+</sup> is administered systemically, there is no damage within the CNS due to the high polarity of the MPP<sup>+</sup> ion which impedes the crossing of the BBB. On the other hand, when MPP<sup>+</sup> is administered intracerebroventricularly (i.c.v.), toxicity in the CNS does occur (Sundström et al. 1986; Giovanni et al. 1994a, b; Sonsalla et al. 2008). Once crossed the BBB, MPTP can interact with MAO-B placed in several cell types (including astrocytes and serotonergic neurons). Since glial cells are abundant in the CNS, MAO-B in astrocytes are thought to be mainly involved in MPTP metabolism (Brooks et al. 1989), while a damage to serotonin neurons fails to protect from MPTP toxicity (Melamed et al. 1986). Once formed MPP<sup>+</sup> is released from glia in the extracellular space, and it enters within DA neurons via the high-affinity DA transporter (DAT, Javitch et al. 1984) (Fig. 2). This was first demonstrated by preventing MPTP-induced DA toxicity using selective DAT inhibitors (Pileblad and Carlsson 1985; Ricaurte et al. 1985; Sundstrom & Jonsson 1985). Uptake selectivity contributes to the selective toxicity of MPP<sup>+</sup>. In fact, MPP<sup>+</sup> is also taken up by norepinephrine (NE) neurons through the NE transporter (NET). Indeed, the uptake within NE neurons exceeds that occurring in DA cells in spite of the fact that NE neurons were originally considered not to be involved in MPP<sup>+</sup> toxicity (Herkenham et al. 1991). Now we are aware that the dosing protocol of MPTP is critical in producing a damage which also involves NE neurons (Seniuk et al. 1990). Remarkably, this point provides a further convergence between MPTP and PD since the damage to NE neurons mostly placed in the nucleus locus coeruleus (LC) is a typical feature of PD patients (Gesi et al. 2000; Zarow et al. 2003, see also in this chapter the Sect. 8). Once entered the DA/NE axons, MPP<sup>+</sup> is accumulated in the cytoplasm where it forms a complex with neuromelanin (D'Amato et al. 1987), being otherwise stored within DA/NE vesicles through the vesicular monoamine transporter type 2 (VMAT-2) (Fig. 2). From these storage sites, MPP<sup>+</sup> is released into the cytoplasm where it may enter the mitochondrion (Ramsay and Singer 1986), which is believed to be the ultimate target to produce neurotoxicity (Fig. 2). Therefore, it is expected that reducing the entry of MPP<sup>+</sup> into

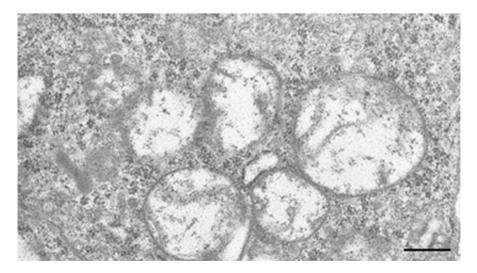


**Fig. 2** Kinetics and dynamics of MPTP in the CNS. Once injected MPTP, which is a lipophilic compound, easily crosses the blood–brain barrier. This allows MPTP to enter within the CNS, where it is converted by monoamine oxidase (MAO)-B into MPP<sup>+</sup> which is actively secreted from glia in the extracellular space. MPP<sup>+</sup> is taken up quite selectively mostly by catecholamine neurons through the high-affinity DA transporter (DAT as shown in the cartoon) by also the NE transporter (NET, not shown). This allows MPP<sup>+</sup> to access DA axons where it can be stored either within synaptic vesicles through the vesicular monoamine transporter (VMAT) or within mitochondria. MPP<sup>+</sup> within mitochondria binds to complex I to inhibit NADH dehydrogenase. The production of reactive oxygen species (ROS) and other free radicals may be the consequence of mitochondrial impairment and the direct oxidative effects of MPP<sup>+</sup> on specific proteins and the oxidative effects of DA which is released by MPP<sup>+</sup> from synaptic vesicles on specific targets. Increased amount of oxidative species produced by all these mechanisms is expected to alter the folding of a variety of cell proteins. This leads to the accumulation of misfolded proteins which mostly consist of co-chaperones such as alpha-synuclein. The amount of oxidative stress is increased by the concomitant opening of calcium ( $Ca^{2+}$ ) channels which may derive from the overactivation of glutamate receptors

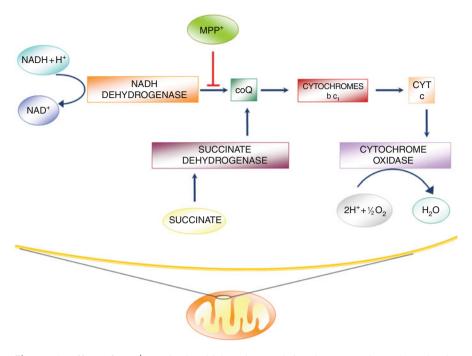
the axons or increasing the storage into the vesicles should protect from MPTP toxicity (Staal and Sonsalla 2000). In fact, MPTP toxicity does not occur in DAT knockout (KO) mice (Gainetdinov et al. 1997), while MPTP toxicity is increased in VMAT-2 heterozygote KO mice (Gainetdinov et al. 1998). In line with this, mice with a 50 % depletion of VMAT show increased vulnerability to MPTP (Takahashi et al. 1997). Thus, synaptic vesicles represent storage sites which buffer MPP<sup>+</sup> and prevent its interaction with mitochondria and other cytosolic targets of toxicity (i.e., chaperone proteins). Intracellular compartmentalization of MPP<sup>+</sup> is critical to understand species differences in MPTP toxicity (Fig. 2).

# **3** MPP<sup>+</sup> and Mitochondria

Although the ultimate mechanisms of cell death induced by MPP<sup>+</sup> remain unclear, strong evidence indicates that MPP<sup>+</sup> inhibits mitochondrial function. MPP<sup>+</sup> enters into mitochondria to inhibit NADH dehydrogenase within mitochondrial respiratory chain (complex I, Fig. 2, Nicklas et al. 1985; Ramsay et al. 1986), thus binding to the same site as rotenone, which is a classic complex I inhibitor (Heikkila et al. 1985; Ramsay and Singer 1986; Krueger et al. 1990). This leads to morphological damage of mitochondria as shown in Fig. 3. As shown in Fig. 4, impairment of mitochondrial respiration by MPP<sup>+</sup> is believed to decrease the levels of oxidized nicotinamide adenine dinucleotide (NAD) (Sonsalla et al. 1992). In line with this,  $MPP^+$  neurotoxicity is related to the failure of energy supplies in a number of in vitro and in vivo experimental settings (Di Monte et al. 1986; Denton and Howard 1987). For instance, within synaptosome preparations, MPP<sup>+</sup> dose-dependently decreases adenosine triphosphate (ATP) content (Scotchner et al. 1990). These effects produce a failure of electrogenic pumps such as 3Na<sup>+</sup>-2K<sup>+</sup> ATPases (Storey et al. 1992), which determines depolarization which leads to a sudden release of neurotransmitters. In fact, acute DA depletion caused by MPTP is preceded by a decrease in ATP levels in vivo within mouse striatum (Chan et al. 1993a). Consistently, the depletion of energy stores occurs more in those regions which take up selectively MPP<sup>+</sup> such as the nigrostriatal DA system (and other catecholamine-containing areas both in the CNS and periphery). This is



**Fig. 3** Electron micrograph of mitochondria following MPTP exposure. MPTP exposure targets mitochondria which are functionally impaired but also deranged in their fine structure. This electron micrograph shows mitochondria within DA neurons from a C57 black mouse treated with MPTP (20 mg/Kg, X4, 2 h apart). Mitochondria lose their crests and undergo severe matrix dilution. Bar=0.24µm (Unpublished data, courtesy of Dr. P Lenzi, University of Pisa)



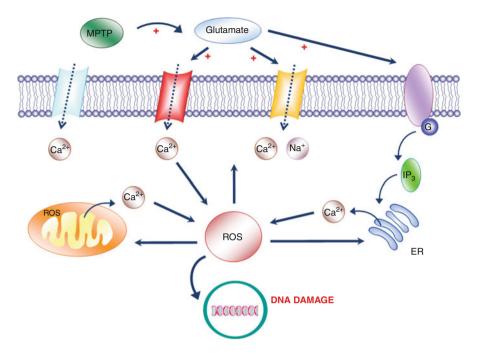
**Fig. 4** The effects of MPP<sup>+</sup> on mitochondrial respiratory chain. The cartoon shows the molecular target of MPP<sup>+</sup> within the mitochondrial respiratory chain at the level of complex I. This corresponds mainly to the same site targeted by rotenone (see " $\triangleright$  Rotenone as Preclinical Model Compound in Parkinson Disease" in this book) and it affects the enzyme NADH dehydrogenase. Thus, MPP<sup>+</sup> blocks NADH dehydrogenase, which produces a variety of downstream effects consisting of ATP loss, formation of ROS, and induction of apoptotic proteins

why, apart from DA toxicity, even the transient DA depletion, which occurs suddenly following MPTP administration, can be prevented by DA uptake inhibitors (Chan et al. 1991).

The identification of mitochondria as the primary target of MPTP-induced toxicity cops well with mitochondrial defects occurring in idiopathic parkinsonian patients as well as the mutations in mitochondrial proteins described in genetic forms of PD (Gu et al. 1998; Dodson & Guo 2007; Büeler 2009).

# 4 MPP<sup>+</sup> and Free Radicals

In physiological conditions free radicals, including reactive oxygen species (ROS), are constantly produced as the consequence of several chemical reactions in the cell including mitochondrial metabolism (Figs. 2 and 4). Routinely, this is balanced by antioxidants, free radical scavengers, and a variety of chaperones, which lead to either innocuous end-products or detrimental compounds. Antioxidants include



**Fig. 5** Glutamate binds to ionotropic and metabotropic receptors. MPTP exposure increases the level of extracellular amino acid transmitters. In the case of this cartoon, glutamate overactivity stimulates both ionotropic (NMDA, kainate and AMPA) receptors and several subtypes of metabotropic receptors. This glutamate overactivity leads to the entry of a variety of ions. In particular, calcium entry, which is also modulated by metabotropic receptors acting on IP<sub>3</sub>, is a strong candidate to contribute to free radical and ROS increase in DA neurons

superoxide dismutase (SOD), glutathione peroxidase, catalase, glutathione (GSH) and coenzyme Q10, essential nutrients (vitamin C, vitamin E, selenium, etc.), and dietary compounds (such as bioflavonoids). Oxidative stress results from an imbalance between formation and neutralization of pro-oxidants.

Mitochondrial impairment increases the production of free radicals (Fig. 2), and these effects may be enhanced by glutamate overactivity (Fig. 5; see also the Sect. 6, in this chapter), being one the consequence of the other. Impairment of mitochondrial function and decrease of cellular ATP may increase free radicals and mostly ROS (Cleeter et al. 1992).

Mitochondrial ROS react with several targets, including manganese SOD (Mn-SOD, SOD2), nitric oxide (NO), and susceptible iron–sulfide (Fe–S)-containing proteins. Mn-SOD is an antioxidant enzyme which protects against ROS produced by mitochondria. In fact, overexpression of Mn-SOD in transgenic mice protects against MPP<sup>+</sup> toxicity (Klivenyi et al. 1998; Maragos et al. 2000). On the other hand, heterozygous Mn-SOD KO mice show higher DA depletion and significantly larger striatal lesions after systemic MPTP treatment compared with controls (Andreassen et al. 2001). Again, overexpression of copper–zinc SOD

(Cu–Zn SOD, SOD1) in transgenic mice protects against MPTP neurotoxicity (Przedborski et al. 1992), and similar effects occur in mice overexpressing either human wild-type SOD1 or human mutant SOD1-G93A (Fornai et al. 2002). This may be due to the interaction of SOD1 with mitochondrial complex I activity thus competing with MPP<sup>+</sup> (Browne et al. 1998; Fornai et al. 2002).

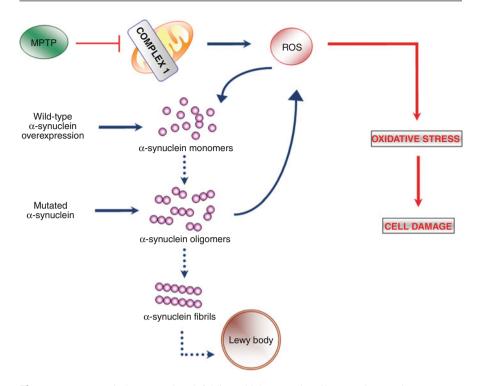
It is puzzling that, apart from conferring protection against MPTP toxicity, the mutation in the gene coding for the human SOD1 leads to a spontaneous loss of nigral DA neurons (Andreassen et al. 2001).

# 5 MPTP, Protein Clearing Mechanisms, and Apoptosis

When acutely administered, MPTP does not affect much protein clearance, but directly targets the mitochondria. In contrast, when MPTP intoxication is continuously protracted for a few weeks, it impairs protein clearing mechanisms (Fornai et al. 2005). This is due mainly to the oxidation of specific proteins such as alpha-synuclein which works as co-chaperone (Fig. 6, Chandra et al. 2004, 2005). In fact following MPTP administration, there is an increased expression of alpha-synuclein in the cytoplasm (Fornai et al. 2005; Purisai et al. 2005) which may induce the formation of toxic alpha-synuclein oligomers (Stefanis et al. 2001; Tanaka et al. 2001; Bucciantini et al. 2002; Walsh et al. 2002; Giorgi et al. 2006), thus leading to neuronal toxicity (Fig. 6).

In this way, despite of its role as co-chaperone (Chandra et al. 2005), alphasynuclein may become a detrimental compound for the survival of both DA and NE neurons once oxidized in excess during MPTP intoxication (Figs. 6 and 7, Conway et al. 2000; Fornai et al. 2005). In fact, chronic or continuous exposure to MPTP (Fornai et al. 2005; Meredith et al. 2008) is accompanied by pathological protein aggregates containing ubiquitin and alpha-synuclein since free radicals and ROS react with alpha-synuclein producing toxic oligomers which cannot be metabolized (Conway et al. 2000; Sulzer 2001) (Figs. 6, 7 and 8).

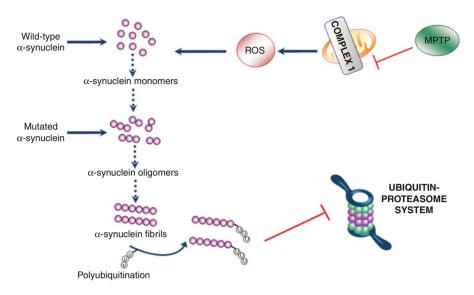
In order to counteract the formation of detrimental protein adducts, the proper activity of protein clearing pathways is critical. This explains why, in recent years, a variety of research studies both on MPTP toxicity and PD focussed on protein clearing systems. The ubiquitin–proteasome (UP) system (Fig. 7) and autophagy (ATG, Fig. 8) are the major systems that degrade intracellular components in eukaryotic cells. The UP system (Fig. 7) mainly serves to degrade short-lived misfolded proteins (Ciechanover 2006). These proteins are targeted by polyubiquitin conjugation, which addresses them to proteolytic degradation within the proteasome core. The ATG pathway is involved in the degradation of long-lived proteins, protein aggregates, and organelles, such as mitochondria, which are delivered to lysosomes for digestion (Fig. 8, Mariño and López-Otín 2004). ATG was found to be increased within the substantia nigra of PD patients by Anglade and collaborators (1997). These authors found the coexistence of ATG vacuoles and apoptosis within melanized neurons of the human substantia nigra. The neurons undergoing an excess of ATG possess condensation of chromatin, moderate vacuolation of endoplasmic



**Fig. 6** MPTP recapitulates protein misfolding which may primarily occur in genetic PD. MPTP administration, mainly via inhibition of mitochondria, or directly, or via DA release, produces high amount of oxidative species which can interact with co-chaperones such as alpha-synuclein to stabilize this protein in oligomeric forms. These oligomers are also produced as a consequence of genetic mutations in alpha-synuclein gene as occurs in a few cases of inherited PD. In all cases alpha-synuclein adducts are toxic for the cells and can either precipitate to form Lewy bodies or spread to specific cell compartments to impair key biochemical pathways thus contributing to DA toxicity

reticulum, and lysosome-like vacuoles containing cytoplasmic material. The existence of dual neuronal death suggests the occurrence of both ATG and apoptosis at different time intervals during degeneration of DAergic cells, with apoptosis being the final stage. This is in line with the recent evidence showing that in DAergic cells dysfunctional ATG triggers apoptosis (Fig. 9, Castino et al. 2008).

A defective protein clearance is supposed to lead to protein accumulation even when misfolded proteins are not produced in great amount. This is in line with the fact that MPTP intoxication, just like PD, produces both ATG alterations and apoptosis (Fig. 9). This is witnessed by the accumulation of altered mitochondria (Fig. 3) and large ATG vacuoles which follows MPTP intoxication. This explains why rapamycin, by activating ATG and promoting lysosomal activity (Fig. 10), attenuates MPTP-induced DA neurodegeneration, thus promoting neuroprotection (Dehay et al. 2010). In addition, in MPTP-treated mice rapamycin has been reported to block the induction of RTP801 gene that promotes neuronal death,

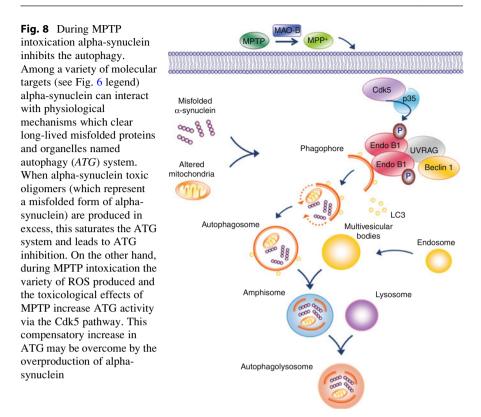


**Fig. 7** During MPTP intoxication alpha-synuclein inhibits the proteasome. Among a variety of molecular targets (see Fig. 6 legend), alpha-synuclein can interact with physiological mechanisms which clear short-lived misfolded proteins named ubiquitin–proteasome (UP) system. When alpha-synuclein toxic oligomers (which represent a misfolded form of alpha-synuclein) are produced in excess, this saturates the UP system and leads to UP inhibition

which is highly expressed in DA neurons of the substantia nigra both in MPTP-treated mice and PD patients (Malagelada et al. 2006).

Remarkably, Liu et al. (2013) found that rapamycin rescues the number of DA neurons and reduces the increase of alpha-synuclein immunoreactivity induced by MPTP. These studies demonstrate that rapamycin via activation of ATG/lysosome pathways both prevents and reverts MPTP toxicity.

Thus, most evidence indicates that MPTP-induced neuronal death is the consequence of energy failure and oxidative stress that occur early on after MPTP administration. In turn, these phenomena are pivotal in triggering molecular pathways which, once activated, rapidly lead to the death of intoxicated neurons (Przedborski and Vila 2003). In fact, the removal of damaged mitochondria and misfolded proteins occurring following MPTP administration is critical in modulating MPTP toxicity. In fact, the occurrence of misfolded proteins in neuronal bodies following prolonged MPTP administration is accompanied by the impairment of the UP as a protein clearing system (Fig. 7, Fornai et al. 2005). Similarly, defective clearance of damaged mitochondria and long-lived proteins cleared by ATG is the hallmark of cell pathology in PD and following MPTP as well (Fig. 8). The abundance of altered mitochondria and toxic protein species is pivotal in triggering apoptotic cell death (Fig. 9). This is in line with the concomitant occurrence of altered ATG and apoptosis both in PD patients (Anglade et al. 1997) and following MPTP intoxication (Novikova et al. 2006). Thus, MPTP can induce apoptosis by different mechanisms. In fact oxidative stress and impairment

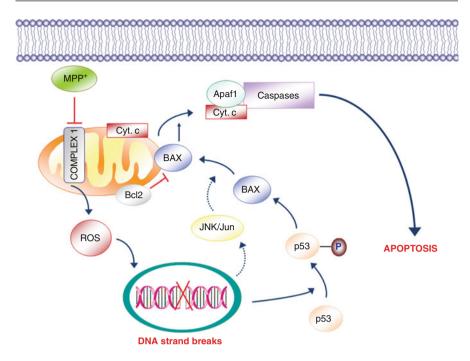


of mitochondrial respiration, following MPTP treatment, can initiate apoptosis (Hartley et al. 1994), and MPTP also induces mitochondrial permeability transition (MPT), which determines mitochondrial swelling and cytochrome c release, with subsequent caspase-dependent apoptosis (Cassarino et al. 1999; Lee et al. 2006). MPTP-induced apoptosis involves different molecules and pathways, including DNA strand breaks, the c-Jun N-terminal kinase (JNK) pathway, caspases, Par-4 protein, p53, Apaf-1, and Bax (Eberhardt and Schulz 2003) (Fig. 9).

# 6 MPTP and Glutamate

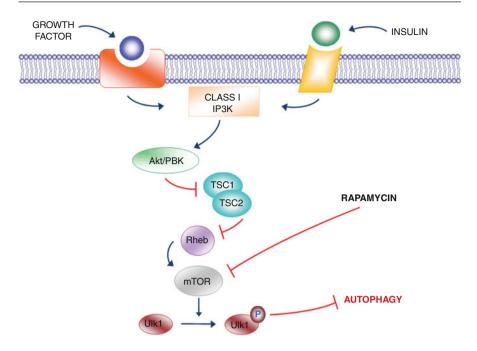
Other sources of free radicals during MPP<sup>+</sup> toxicity besides mitochondria are glutamate and DA itself. Robust experimental evidence indicates that glutamate and other excitatory amino acids (EAA) play a role in MPTP toxicity since chronic MPTP increases extracellular glutamate in the substantia nigra (Meredith et al. 2009) and MPP<sup>+</sup> increases extracellular levels of glutamate in the striatum (Carboni et al. 1990).

The acidic amino acids glutamate, aspartate, and their analogues bind to three main types of ionotropic and also to metabotropic receptors (Fig. 5). These amino acids are well known to be responsible for "excitotoxicity" (Olney et al. 1980),



**Fig. 9** The effects of MPP<sup>+</sup> on mitochondria trigger apoptotic proteins. The cartoon shows the molecular target of MPP<sup>+</sup> within the mitochondrial respiratory chain at the level of complex I. This inhibition leads to production of a variety of downstream effects consisting of ROS formation and induction of apoptotic proteins such as BAX and Bcl-2

which indicates how endogenous glutamate overactivity becomes frankly toxicant. These conditions include excessive and/or prolonged release of glutamate into the synaptic cleft. In these cases, neuronal loss progresses downstream along synaptically joined neurons, both after repeated acute insults and in neurodegenerative diseases in which the "systemic degeneration" consists of spreading cell loss to neurons interconnected within functional circuits. This might be the case of glutamatergic pathways from the neocortex to the SNpc and striatum (Cotman et al. 1987). In keeping with these concepts, the study of toxicity progression along the circuitries enrolled during acute insults (MPTP) provided the pathophysiological basis to depict the regional impairment occurring also in neurodegenerative diseases (PD). In fact the nigrostriatal pathway receives synaptic terminals from glutamatergic neurons and possesses EAA receptors. Furthermore interrupting the excitatory circuit by lesioning upstream neurons or trans-sectioning the glutamatergic input to the nigrostriatal pathway attenuates the toxic insult. This is in line with the fact that a noncompetitive N-methyl-D-aspartate (NMDA) antagonist MK-801 (Chan et al. 1993b) protects against MPTP toxicity in rodents (Turski et al. 1991; Storey et al. 1992) and primates (Zuddas et al. 1992). An



**Fig. 10** Effects of rapamycin on autophagy. The ATG pathway can be modulated by a number of compounds among which growth factors and insulin are shown in the cartoon. Independently by the way of inducing ATG, rapamycin is a strong ATG inducer which promotes ATG downstream in the regulatory chain at the level of mTOR which is a strong ATG inhibitor. Rapamycin, by inhibiting mTOR, powerfully activates ATG

analogous protection was obtained by damaging glutamatergic corticostriatal fibers (Storey et al. 1992) suggesting that endogenous excitotoxicity contributes to MPTP toxicity (Brouillet and Beal 1993; Lange et al. 1993; Srivastava et al. 1993; Lange and Riederer 1994; Loschmann et al. 1994; Ossowska 1994; Vaglini et al. 1994; Kanthasamy et al. 1997; Sonsalla et al. 1998; Araki et al. 2001).

Species differences play a major role in the effects of glutamate in MPTP toxicity (Fornai et al. 1997b) leaving under debate the relevance of this point for PD patients.

As mentioned above, some authors have shown an involvement of EAA in the toxicity of MPP<sup>+</sup> injected directly into the substantia nigra (Turski et al. 1991) or the striatum (Storey et al. 1992) which was counteracted by the simultaneous administration of a competitive or noncompetitive NMDA receptor antagonist to the substantia nigra. Other authors could not confirm these results (Sonsalla et al. 1992) nor did they observe any involvement of EAA after systemic injection of MPTP in mice (Sonsalla et al. 1989, 1992; Kupsch et al. 1992). The way of administering MPTP also appears to be crucial. This concept is in line with the results of Storey et al. (1992), who showed that MPP<sup>+</sup> administered within the rat striatum produces long-lasting effects on neuronal energy metabolism, thus causing a three- to fourfold decrease in ATP levels, which persists for over 48 h.

Although rodent data are contradictory, studies performed on MPTP-treated primates, both old- and new-world monkeys: (Zuddas et al. 1992; Lange et al. 1993), respectively, demonstrated a clear protective role of NMDA and non-NMDA ionotropic receptor antagonists (both competitive and noncompetitive) in MPTP-induced parkinsonism (Crossman et al. 1989; Close et al. 1990).

The role of glutamate is not limited to its effects on ionotropic receptors and involves the role of metabotropic receptors which might be a therapeutic target with less side effects. In detail it was demonstrated that mGlu5 receptor agonists exacerbate, while antagonists may counteract MPTP-induced nigrostriatal damage (Battaglia et al. 2004). Nowadays it seems likely that glutamate per se is not the mediator of MPTP toxicity. The neurotoxic mechanisms rather rely on other pathways, and this brings the role of glutamate in the sideshow along the mainstream leading to MPTP toxicity.

# 7 MPTP and Dopamine

As reported, MPP<sup>+</sup> provokes a sudden, quite generalized neurotransmitter release. This phenomenon is more intense for catecholamine-containing neurons including DA cells. At this level, MPP<sup>+</sup> induces massive DA release from storage vesicles into the cytosol, where DA itself produces free radicals via auto-oxidation or MAO-mediated oxidative deamination, resulting in the formation of hydrogen peroxide, superoxide, and reactive quinone species (Lotharius and O'Malley 2000). The latter are known as DA quinones which also derive by activating cyclooxygenase (COX)-2 (Teismann et al. 2003). This may explain the neuroprotective effects of COX-2 inhibitors against MPTP toxicity. In addition, the activity of tyrosine hydroxylase (TH) may produce ROS as a normal by-product.

To emphasize the detrimental effects induced by DA within DA neurons in experimental parkinsonism induced by MPTP, it is useful to mention the toxic mechanisms induced by methamphetamine (METH), a neurotoxic agent which releases DA even more powerfully than MPTP (refer to specific chapters on ▶ Methamphetamine and MDMA Neurotoxicity: Biochemical and Molecular Mechanisms, this volume). Administration of METH produces experimental parkinsonism by inducing the loss of nigrostriatal DA axons (Ricaurte et al. 1982; Woolverton et al. 1989; Ricaurte and McCann 1992; McCann et al. 1998; Villemagne et al. 1998; Volkow et al. 2001a, b; Battaglia et al. 2002) and DA perikarya (Sonsalla et al. 1996; Castino et al. 2008; Granado et al. 2011). As for MPTP, METH abuse was recently shown to facilitate PD (Callaghan et al. 2010, 2012). As MPTP, METH is transported into DA terminals where it interacts with DAT and VMAT-2. In this way, METH causes massive DA release (O'Dell et al. 1991; Cubells et al. 1994). This reference to METH is key in the context of understanding the toxic potential of DA-releasing neurotoxicants. In fact, a high DA release occurs also during MPTP intoxication. High DA levels promote DA oxidation which is toxic to a variety of key proteins and organelles within DA neurons (Castino et al. 2008). The role of DA toxicity can lead to an understanding of some critical questions about MPTP such as species differences. In fact, rats are resistant to MPTP toxicity, while mice are more sensitive. When looking at the kinetics and dynamics of MPTP in mice and rats, no difference between mice and rats appears but a remarkable discrepancy in MPTP-induced DA release. Mice, which are sensitive to MPTP toxicity, undergo massive MPTP-induced DA release, while rats which are almost resistant to MPTP toxicity undergo a minimal DA release following MPTP administration. This concept is extensively reviewed in the paragraph titled "Species Differences to MPTP Toxicity" but is useful to be mentioned here to understand the role of DA in MPTP toxicity. It is quite intriguing that DA as a neurotransmitter possesses such a great potential to produce neurotoxicity. This was already demonstrated by pioneer studies injecting DA directly in the striatum and reviewing the potential toxicity which follows DA release. The ability of DA to act as a neurotoxicant relates to the chemical reactivity reviewed so far. In contrast, NE does not produce toxicity but it is rather neuroprotectant against MPTP toxicity (Mavridis et al. 1991; Marien et al. 1993; Fornai et al. 1995, 1996; 1997a).

# 8 MPTP Toxicity and Norepinephrine

Damage to the NE nucleus LC is known to worsen the onset and progression of experimental parkinsonism induced by MPTP both in rodents and monkeys (Mavridis et al. 1991; Marien et al. 1993; Bing et al. 1994; Fornai et al. 1995; Gesi et al. 2000). Damage to NE axon terminals, induced by the selective NE neurotoxin N-(-2-chloroethyl)-N-ethyl-2-bromobenzylamine (DSP-4) (Fornai et al. 1996) before MPTP administration, increases the sudden MPTP-induced DA release and enhances MPTP toxicity (Fornai et al. 1997a). The role of NE in MPTP toxicity directly applies to human PD patients where analogous effects were described (Rye and DeLong 2003).

In fact, the loss of NE is thought to facilitate the onset and progression of PD (Tong et al. 2006; Rye and DeLong 2003). Actually, the loss of LC neurons also represents a hallmark of PD since extensive neuropathological analysis demonstrated that LC neurons are lost even more than SNpc cells in PD (Zarow et al. 2003). The loss of LC neurons in PD represents a pioneer finding dating back at the beginning of the last century (Tretiakoff 1919). It remained a smoldering concept up to experimental studies using MPTP and showing a causal link between LC damage and the loss of DA-containing neurons. Mavridis et al. (1991) developed a primate model demonstrating that previous damage to LC NE-containing neurons may impair the recovery of nigrostriatal DA pathway following MPTP intoxication. Further studies demonstrated that a loss of NE innervation causes an increased susceptibility of DA neurons to MPTP (Fornai et al. 1997a). It is worth to be mentioned that such a reduced threshold within DA neurons to MPTP neurotoxicity is not due to a prolonged persistence of MPP<sup>+</sup> within the CNS. In fact, since MPP<sup>+</sup> is taken up within NE neurons at a rate which is faster compared with DA neurons (Herkenham et al. 1991), it is expected that a damage to NE cells would have produced a loss of what may be considered a sort of buffering activity. Paradoxically, LC-damaged brains retain MPP<sup>+</sup> for a longer time compared with LC-intact

brain. This indicates that variations in toxin kinetics are not causing enhancement of MPTP toxicity in NE-damaged brains (Fornai et al. 1997a). It is rather the occurrence of NE or the absence of such a neurotransmitter which occludes or sensitizes, respectively, MPTP-induced DA toxicity. Enhancement of MPTP toxicity occurs also by preventing NE synthesis using a DA beta-hydroxylase inhibitor such as fusaric acid or disulfiram (Corsini et al. 1985; Weinshenker et al. 2008). Thus, the protective effects of NE-containing neurons on MPTP toxicity are demonstrated throughout a variety of experimental approaches and animal species (Fornai et al. 2007a). These findings pose the need for well-preserved LC neurons in order to prevent the occurrence of PD. In fact, in human patients Tong et al. (2006) demonstrated a tight correlation between early loss of NE- and the later damage to DA-containing neurons within specific brain areas of PD patients. These authors suggested that a loss of NE innervations anticipates subsequent damage to DAcontaining cells. On the other hand, if PD is characterized by the concomitant loss of LC NE- and SNpc DA-containing neurons, a neurotoxin which is able to mimic PD is also expected to reproduce such a combined neurotoxicity. This poses the following question: Is there a neurotoxic effect of MPTP for NE neurons? The study of Seniuk et al. (1990) indicates that small reiterated doses of MPTP were effective in damaging NE more than DA neurons (Seniuk et al. 1990). Pathobiochemical (Hornykiewicz 1975, 1998; Tong et al. 2006) and pathological findings (Braak et al. 2000; Zarow et al. 2003) clearly established the concomitant NE and DA damage in PD. A number of studies on MPTP toxicity up to the recent findings in mice (Meredith et al. 2002; Fornai et al. 2005) and nonhuman primates by Masilamoni et al. (2011) confirm that prolonged exposure to MPTP does produce a concomitant damage to DA and NE neurons as demonstrated in PD.

These findings introduce the issue of whether we should consider prolonged MPTP exposure more reliable than acute administration to produce toxic effects which mimic PD. This point extends to the evaluation of various protocols of MPTP administration which were developed during the last decades.

### 9 Acute, Chronic, and Continuous MPTP Administration

In mice a wide range of MPTP administrations were experimented. These vary with respect to the total amount of MPTP administered, the route (intraperitoneally versus subcutaneously), and time between injections, including the acute (usually four doses within 6 h), the subacute (usually 5 days), and the chronic administration (weeks), the latter often accompanied by an adjuvant drug, probenecid (Petroske et al. 2001; Kurosaki et al. 2004; Lau et al. 2005; Anderson et al. 2006). Very recently (Prediger et al. 2011) mice have been treated with intranasal MPTP, showing impairments in olfactory, cognitive, emotional, and motor functions, similarly to those observed in PD. This was accompanied by TH loss in the olfactory bulb and SNpc, with significant DA depletion in different brain areas (Prediger et al. 2011).

The different MPTP administrations result in different amount of DA depletion, motor deficits, and time course of recovery (Sonsalla and Heikkila 1986;

Jackson-Lewis et al. 1995; Bezard et al. 1997a, b; Kühn et al. 2003). When injected into mice, MPTP causes a PD-like syndrome with massive loss of nigral DAergic neurons and striatal DA. Acute administration of MPTP leads to striatal loss that can range from 40 % (when MPTP is given at 14 mg/kg per dose four times a day) to approximately 90 % (20 mg/kg per dose four times, 2 h apart in a single day) 7 days after the last MPTP dose and at least 40-50 % loss of nigrostriatal DAergic neurons in young adult C57BL/6 mice by 7 days after MPTP administration (Jackson-Lewis and Przedborski 2007). A subchronic regimen consisting of administration of systemic MPTP 30 mg/kg per day for 5 consecutive days in Swiss mice leads to morphologic features resembling apoptotic cell death occurring on day 6 after the last MPTP administration, with tissue DA depletion of 59 % 1 day after MPTP last administration (Tatton and Kish 1997) and a progressive recovery of DA depletion (about 28 %) from day 3 to day 14 after MPTP discontinuation (Serra et al. 2002). During the acute MPTP administration, necrotic rather than apoptotic features of the neurons of the SNpc were observed (Jackson-Lewis and Przedborski 2007); this suggests that the schedule of MPTP administration does not only influence the time course of nigrostriatal damage but may also provide insights into the underlying mechanisms of PD pathogenesis, such as the induction and manifestation of neuronal death (necrotic or apoptotic), which appears to be correlated to different stages of the human PD (presymptomatic, immediate onset, progressive, and final stage).

Shall we expect that, in order to better reproduce PD, MPTP needs to be administered once in high doses or rather being reiterated in small amount during long time intervals or even administered continuously?

In this context when considering the effects of acute MPTP administration to a variety of animal species, it is evident that despite being able to reproduce a massive neuronal loss in the SNpc, such an experimental protocol leaves NE neurons intact and does not produce the occurrence of neuronal inclusions neither in mice nor in primates. In fact, although intraneural inclusions containing eosinophilic materials and alpha-synuclein aggregates have been described in MPTP-treated monkeys (Forno et al. 1986; Kowall et al. 2000), the classic Lewy bodies, a typical feature of PD, have not been convincingly observed (Forno et al. 1993). The intraneural inclusions containing eosinophilic material and alpha-synuclein that are reminiscent of Lewy bodies have not been observed in young monkeys (Forno et al. 1995). In fact, following acute administrations the time course of MPTP-induced neurotoxicity is rapid, and therefore it does not activate long-acting neuronal mechanisms which are required to build neuronal bodies. In fact, it is likely that neuronal inclusions represent a compromise between long-acting slight neurotoxic mechanisms and the recruitment of compensatory mechanisms. This is what is thought to happen during PD which possesses a chronic progressive disease course.

On the other hand, when MPTP is administered chronically or continuously in mice apart from NE damage, also neuronal bodies can be described (Meredith et al. 2008; Fornai et al. 2005; Gibrat et al. 2009; Beal 2011). Sometimes chronic models based on repetitive small doses remain subthreshold to induce toxicity as reported in one study (Alvarez-Fischer et al. 2008), while most studies found that chronic

(Meredith et al. 2008) or continuous (Fornai et al. 2005; Cleren et al. 2008; Gibrat et al. 2009; Yasuda et al. 2011) MPTP administration protracted for a few weeks does produce alpha-synuclein-positive neuronal bodies in addition to massive striatal DA depletion and SNpc neuronal loss. The amount of MPTP and the time needed to produce inclusions may vary between different studies; this is likely to depend on experimental variability such as purchaser and housing which are well known to modulate the threshold for MPTP toxicity (Heikkila et al. 1984b; see also the Sect. 14).

Prolonged neurotoxin administration is effective also for the active metabolite MPP<sup>+</sup> when this is delivered continuously across the blood–brain barrier both in rodents and primates (Imai et al. 1988; Sirinathsinghji et al. 1988). In the rat, continuous *i.c.v* MPP<sup>+</sup> administration produces striatal DA loss and SNpc cell death as well as neuronal bodies staining for ubiquitin and alpha-synuclein (Yazdani et al. 2006).

### 10 Spinal Cord and MPTP

The classic pathological features of sporadic PD include the degeneration of DA neurons in the SNpc and NE neurons in the LC. Besides these areas, also the spinal cord (SC) seems to be affected in PD in the presymptomatic stage. This was already demonstrated by significant motor neuron loss in various PD models (Bloch et al. 2006; Klos et al. 2006; Samantaray et al. 2008a), including the A53T mutant mice that express the mutant alpha-synuclein gene of familial PD (Martin et al. 2006). Following MPTP administration in the SC, specific apoptotic proteins are highly expressed (Chera et al. 2002, 2004; Samantaray et al. 2006, 2007). In a very recent paper, it was found that in MPTP-treated mice there is a significant motor neuron loss in the lumbar and cervical SC (Vivacqua et al. 2012). Samantaray and colleagues (2008b) showed that MAO-B are expressed in the SC and DAT is expressed in motor neurons thus suggesting that MPTP can be converted to MPP<sup>+</sup> which in turn might be taken up by motor neurons to induce cell death.

# 11 MPTP, the Heart, and the Gut

While MPTP toxicity in the CNS is extensively investigated, peripheral autonomic dysfunction following MPTP intoxication received much less attention. Only recent studies demonstrated a parallelism between PD and MPTP in producing cardiac denervation due to a loss of NE axons in the heart (the so-called Parkinsonian heart; Fukumitsu et al. 2006; Fornai et al. 2007b). Similarly, MPTP intoxication was recently reported to reproduce the decrease in plasma testosterone levels observed in PD, and this was related with alteration produced by MPTP in Leydig cells (Ruffoli et al. 2008). Gastrointestinal (GI) dysfunction is a well-recognized feature of PD (Natale et al. 2008) and the entire GI tract may be affected (Edwards et al. 1991; Jost 1997; Pfeiffer 2003). A few studies suggest that GI symptoms in PD may be caused by a neurodegenerative process in the enteric

nervous system (ENS), similar to what occurs in the CNS (Singaram et al. 1995; Wakabayashi et al. 1988), and it is hypothesized that degeneration in the ENS mirrors and precedes analogous pathological events in the brain or SC (Braak et al. 2003, 2006, 2007; Natale et al. 2011).

Only a few experimental studies using MPTP addressed whether such intoxication involves the gut. In rats exposed to MPTP, Szabo et al. (1985) described inhibition of duodenal spike activity. Decreased DA levels were found in the jejunal myenteric plexus, although this occurred only for high doses (Szabo et al. 1985; Eaker et al. 1987).

The effects of MPTP in the mouse colon have been evaluated in vivo, following a total dose of 60 mg/kg of MPTP. This produced a 40 % reduction of TH-positive neurons (Anderson et al. 2007). Whether the loss of TH-positive neurons following MPTP administration was due to DA or NA or both was recently demonstrated by Natale and collaborators (2010). This appears to be due to the selective loss of enteric DAergic neurons, as confirmed by the selective damage to DAT-containing cells and the specific loss of DA in the homogenates from the gut. In the same studies, it was found that MPTP increases alpha-synuclein thus reproducing alterations occurring in the digestive system of PD patients.

# 12 Species Differences in MPTP Toxicity

A variety of animal species are sensitive to MPTP toxicity including monkeys, rodents, cats, pigs, dogs, sheep, rats, and goldfishes (Kopin and Markey 1988; Gerlach and Riederer 1996; Przedborski et al. 2001); however, different effects are produced by MPTP in different species ranging from massive (primates) to minimal (rats) toxicity.

In monkeys, MPTP administration reproduces most of the clinical and pathological hallmarks of PD (Langston et al. 1984a; Chiueh et al. 1984; Doudet et al. 1985; Crossman et al. 1985; Nomoto et al. 1985; Vezoli et al. 2011) but not in rats, which were found to be resistant to MPTP (Chiueh et al. 1984). In detail, rats can be damaged by MPTP only in the periphery, being necessary to administer MPP<sup>+</sup> at very high doses intracerebroventricularly to produce a damage within CNS to DA neurons (Yazdani et al. 2006). Mice are sensitive to MPTP but require higher amount than primates. In detail, in mice MPTP needs to be administered acutely at doses ranging from 15 to 80 mg/Kg to produce a progressively massive loss of nigral DA neurons, while in monkeys the same effects are induced at doses at least 100-fold lower. Such a discrepancy is not related with species differences in the bioavailability of MPP<sup>+</sup> within the striatum following systemic MPTP (Zuddas et al. 1994; Vaglini et al. 1996); it is likely that species differences in the toxic effects of comparable amounts of nigrostriatal MPP<sup>+</sup> may relate to species differences in the pharmacodynamics of MPTP. In fact, it is well known that comparable amounts of MPP<sup>+</sup> are formed within the striatum of mice and rats following similar doses of systemic MPTP (Zuddas et al. 1994; Vaglini et al. 1996). This rules out the chance that MAO-B works differently in these species to convert MPTP into MPP<sup>+</sup>. However, if one considers that MPP<sup>+</sup> is selectively taken up by DA terminals, this process could differ significantly. In a careful series of experiments, Giovanni et al. (1994a, b) found no species difference in the uptake of MPP<sup>+</sup>, ruling out also the difference in DAT efficacy. On the other hand, MPP<sup>+</sup> apart from damaging the mitochondrial respiratory chain is also able to produce a robust DA release (see Sect. 7). It was shown that equimolar amount of MPP<sup>+</sup> produces a pronounced DA release in mice versus a modest effect in rats (Giovanni et al. 1994a, b). This point which remained not further investigated sheds a new light on the potential role of intracellular DA joined with the impairment of mitochondrial respiratory chain in sustaining MPTP neurotoxicity. It would be interesting to test the effects of MPP<sup>+</sup> on striatal DA release in primates.

These latter experiments to our knowledge were never carried out. Nonetheless, it is worth to mention that MPTP toxicity is positively related to the amount of DAT (Watanabe et al. 2005) while is negatively related to the amount of VMAT-2 (Chen et al. 2005). In this scenario, we already reported that MPP<sup>+</sup> is sequestered into synaptic vesicles by VMAT-2, and this mechanism appears to protect rat DA neurons from MPP<sup>+</sup> toxicity (Staal and Sonsalla 2000). In fact, in primates MPTP sensitivity is bound to the lower expression of VMAT-2 compared with other species (Chen et al. 2008). Again, if one looks at the site specificity of DA neuron loss in the substantia nigra following MPTP, there is a negative correlation with the expression of VMAT-2 and MPTP-damaged cells. Conversely, the amount of DAT is higher in those subsets of neurons of the substantia nigra which are more vulnerable to MPTP (Haber et al. 1995).

Among the different animal species, mice are commonly used in MPTP experiments. This is due to easy handling, less ethical restrictions, and lower costs compared with monkeys. Nonetheless, mice are significantly less sensitive to MPTP than primates and do not convincingly develop persistent and progressive motor symptoms unless they are treated following sophisticated protocols (Sundström et al. 1990; Przedborski et al. 2001; Fornai et al. 2005).

### 13 MPTP in Nonhuman Primates

MPTP administration to monkeys is considered the best model to reproduce PD due to similarities in neuropathology and behavior. This model was routinely applied in preclinical studies aimed to test pharmacological and surgical treatments of PD. The first MPTP administration in common marmosets was described by Jenner and colleagues (1984), who observed profound akinesia, rigidity of the trunk and limbs, postural abnormalities, loss of vocalization, and, occasionally, whole body tremor (Jenner et al. 1984). MPTP can be administered to monkey using different routes, such as gavage or stereotactical injection. One of the most common consists in the parenteral administration (subcutaneous, intravenous, intra-arterial, intraperitoneal, or intramuscular; Przedborski et al. 2001). Among systemic routes, MPTP is administered through multiple i.p. and and i.m. injections which induce a progressive generalized parkinsonism or through unilateral intracarotid infusion that determines hemiparkinsonism in the injected side. This is critical since bilateral parkinsonism requires L-DOPA to keep the animal alive (Petzinger and Langston 1998). In contrast, unilateral intracarotideal infusions, which determine a chronic model of hemiparkinsonism, enable the monkeys to feed without supporting therapeutics (Bankiewicz et al. 1986; Przedborski et al. 1991).

MPTP can be delivered acutely or chronically, the former being particularly useful in the assessment of treatments aimed at understanding the mechanisms and reducing the amount of L-DOPA-induced dyskinesias (Boraud et al. 2001; Samadi et al. 2008; Johnston et al. 2010; Riahi et al. 2011; Huot et al. 2011; Koprich et al. 2011; Grégoire et al. 2011).

Although both acute and chronic MPTP monkey models are appropriate for testing novel therapies, the chronic model is more appropriate to test neuroprotective drugs.

Even within primates a strong variability to MPTP toxicity exists. For instance old-world monkeys such as macaques are much more sensitive to MPTP than new-world monkeys such as marmosets.

#### 14 MPTP in Mice

In mice, MPTP determines degeneration of DA neurons which again varies depending on the specific mouse strain (Heikkila et al. 1984a). Motor impairment is difficult to be detected in mice although evidence exists showing a correlation between behavioral and pathological findings in MPTP-treated mice (Goldberg et al. 2011). Within mouse strains there is a wide variation in their sensitivity to the MPTP, the C57BL/6 mouse being the most sensitive, whereas CD-1 and BALB/c appear more resistant and Swiss Webster are very resistant to MPTP toxicity (Sonsalla and Heikkila 1988; Muthane et al. 1994; Hamre et al. 1999; Sedelis et al. 2000). Thus, in order to obtain subtotal nigrostriatal damage, it is much better to use C57BL/6 mice.

Sensitivity of mice to MPTP increases with age independently of the strain. A mechanism of strain-dependent sensitivity to MPTP has been hypothesized by Boyd and collaborators (2007), who reported a divergence in the inflammatory response, by examining JNK and COX-2 signaling induction in C57BL/6 J and Swiss Webster mice. A different sensitivity to MPTP within each strain has been reported depending on the purchaser. Thus, Swiss Webster mice from three different commercial suppliers possess different sensitivity to MPTP (Heikkila 1985). Therefore, it is important for investigators to consider both the strain and the vendor used when developing new treatment protocols for MPTP-induced neurotoxicity.

### 15 MPTP in Rats

Rats injected with MPTP doses comparable to those used in mice do not exhibit any significant DA degeneration. The MPTP toxic doses in rats would be incompatible with rat life if injected systemically; therefore, it is necessary to protect against the peripheral effects of MPTP by coadministering a peripheral adrenergic ganglionic

blocker like guanethidine (Giovanni et al. 1994a). Alternatively one might administer MPP<sup>+</sup> at high doses directly within the CNS. The relative resistance of rats to MPTP toxicity was initially attributed to species differences in MPTP metabolism and/or sequestration of MPP<sup>+</sup>. In fact, monkeys have a much longer striatal retention of MPP<sup>+</sup> (half-life about 10 days) compared with mice (half-life about 4 h) (Johannessen 1991). However, Giovanni et al. (1994a, b) found that although the rat is relatively insensitive to MPTP toxicity, the in vitro MPP<sup>+</sup> uptake into DA synaptosomes, the in vivo striatal extracellular MPP<sup>+</sup> concentrations, and the amount of postmortem striatal MPP<sup>+</sup> levels did not correlate with the higher sensitivity to MPTP of mice compared with rats. Unexpectedly, postmortem striatal MPP<sup>+</sup> levels were higher in rats than in mice (Giovanni et al. 1994a, b; Zuddas et al. 1994). Additionally, previous studies showed no difference in mice compared with rats for striatal synaptosomal MAO-B activity and striatal synaptosomal MPP<sup>+</sup> uptake (Giovanni et al. 1991). Zuddas et al. (1994) confirmed the absence of any significant difference between C57 black mouse and various rat strains in striatal MPP<sup>+</sup> kinetics following systemic MPTP administration. Staal and collaborators (2000) confirmed that there was no significant species difference (mice versus rats) in striatal DAT numbers, suggesting a similar accumulation of MPP<sup>+</sup> within the DA neurons in both species. Thus, lower vulnerability to MPP<sup>+</sup> in rats versus mice could not be explained by lower intracellular levels of MPP<sup>+</sup> in rat DA neurons. MPP<sup>+</sup> is actively accumulated into vesicles by VMAT-2 (Del Zompo et al. 1993; Moriyama et al. 1993), and striatal vesicles isolated from rats have a higher density of VMAT-2 compared with mice and a greater ability to sequester MPP<sup>+</sup>, suggesting a larger storage capacity for MPP<sup>+</sup> in rat vesicles (Staal and Sonsalla 2000). In turn this produces lower amount of DA release following MPTP in rats compared with mice. Pretreatment of mice and rats with a VMAT-2 inhibitor, Ro 4-1,284 (10 mg/kg i.p.), produces a fivefold leftward shift in the MPP<sup>+</sup> dose-response curve and a significant lowering of the EC<sub>50</sub> concentration for MPP<sup>+</sup>-induced damage in rats. This provides evidence for a substantial accumulation of MPP<sup>+</sup> in vivo in the rat striatum and explains why enhanced MPP<sup>+</sup> toxicity occurs when blocking VMAT-2. In mice VMAT-2 inhibition fails to enhance MPP<sup>+</sup> toxicity in the striata, suggesting that vesicular sequestration may not be as significant in providing protection in this species as it occurs in rats. Although VMAT-2 inhibition increases MPP<sup>+</sup> toxicity to striatal DA nerve terminals in rats, the potency of MPP<sup>+</sup> toxicity in the striatum of rats in which VMAT-2 is inhibited is still less pronounced than that in mouse striata exposed to MPP<sup>+</sup>. This implies that mechanisms other than the sole MPP<sup>+</sup> sequestration protect against MPP<sup>+</sup> toxicity to DA neurons in different species (Staal and Sonsalla 2000).

### 16 Conclusion

The study of neurotoxicity induced by MPTP is a seminal research field to understand PD. In this chapter we reported the main concepts developed over the last three decades which led us to understand key molecular steps which are pivotal in mediating MPTP toxicity. Remarkably, these molecular mechanisms turned out to be critical also for the survival of DA neurons in idiopathic PD. This is the case of the role played by DAT and VMAT-2 in conditioning the sensitivity to MPTP neurotoxicity and the tendency of DA neurons to undergo progressive degeneration in PD. Similarly, the mitochondria as targets of MPTP toxicity appear similarly affected by selective mutation of genes leading to PD. Again, the fate of mitochondria and the ability to clear these organelles when being dysfunctional are key in the modulation of MPTP toxicity as well as in the course of PD, where certain gene mutations impair the clearance of mitochondria. This is similar for what we discovered about the role of specific misfolded proteins such as alpha-synuclein.

Our perspective on PD is drastically changed in the past three decades, when novel brain areas as well as peripheral sites are increasingly recognized to be affected in PD patients. It is remarkable that progress on MPTP research led to mimic PD even for these additional areas underlying specific additional symptoms. PD is no longer viewed as a highly specific degeneration which involves solely the DA nigrostriatal pathway; similarly MPTP intoxication covers most of the additional features now evident in PD patients. Nowadays, after three decades, we can say that if MPTP per se did not lead to discovery of the environmental compound which causes PD, indeed the study of MPTP did disclose several molecular and cellular pathways which are critical in the genesis of PD. This latter point corresponds fairly to what we enthusiastically expected from MPTP when it was identified as a causal agent of what it remains, a toxic form of environmental parkinsonism.

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# **Cocaine as a Neurotoxin**

# Teresa Cunha-Oliveira, A. Cristina Rego, and Catarina R. Oliveira

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### Abstract

Cocaine is a widely abused psychostimulant drug, with sympathomimetic properties and intense euphoric effects. Cocaine and some of its toxic metabolites cross the blood-brain barrier and induce neurologic impairments, affecting primarily the prefrontal cortex and basal ganglia. In this review, we discuss the mechanisms involved in brain dysfunction induced by cocaine, focusing on pre- and postsynaptic changes in dopaminergic and glutamatergic

T. Cunha-Oliveira (🖂)

A.C. Rego • C.R. Oliveira

e-mail: a.cristina.rego@gmail.com; catarina.n.oliveira@gmail.com

Center for Neuroscience and Cell Biology (CNC), University of Coimbra, Coimbra, Portugal e-mail: teresa.oliveira@gmail.com

Center for Neuroscience and Cell Biology (CNC) and Faculty of Medicine, University of Coimbra, Coimbra, Portugal

neurotransmission, oxidative stress, and mitochondrial dysfunction. Neurotoxic effects of combinations of cocaine with other drugs are also discussed. In summary, cocaine neurotoxicity may underlie brain dysfunction in cocaine and polydrug abusers and may predispose the brain to neurodegeneration.

Keywords

Cocaine • Dopamine • Glutamate • Mitochondria • Neurotoxicity • Oxidative stress

List of Abbreviati	ions
$Ca^{2+}i$	Intracellular Ca <sup>2+</sup> concentration
	Dopamine transporter
DAO	
•	Dopamine quinone
DARPP32	Dopamine- and cAMP-regulated neuronal phosphoprotein
DOPAC	3,4-dihydroxyphenylacetic acid
DOPAL	3,4-dihydroxyphenylacetaldehyde
ERK	Extracellular-signal-regulated kinase
GPx	Glutathione peroxidase
GSH	Reduced glutathione
$H_2O_2$	Hydrogen peroxide
MAO	Monoamine oxidase
MAPKK/MEK	Mitogen-activated protein kinase kinase/extracellular signal-
	regulated kinase kinase
MDA	Malondialdehyde
NMDA	N-methyl-D-aspartate
$O_2^{\bullet-}$	Superoxide anion
юн	Hydroxyl radical
PARP	Poly (ADP-ribose) polymerase
PD	Parkinson's disease
PKA	Protein kinase A
PP1	Protein phosphatase 1
Ras-GRF-1	Ras protein-specific guanine nucleotide-releasing factor 1
SOD	Superoxide dismutase
VMAT	Vesicular monoamine transporter

# 1 Introduction

Cocaine is an alkaloid extracted from the plant *Erythroxylum coca*, cultivated in the South American countries of Bolivia, Colombia, and Peru. This substance is a widely abused psychostimulant drug, with sympathomimetic properties, used recreationally to increase alertness, relieve fatigue, and increase self-confidence and is abused due to its intense euphoric effects. Cocaine is among the most abused

illicit drugs, used annually by 0.3-0.4 % of the world population aged 15–64 years, with higher prevalence in North America (1.6 %), Oceania (1.5–1.9 %) and Western and Central Europe (1.3 %) (United Nations Office on Drugs and Crime 2012).

### 2 Pharmacology of Cocaine

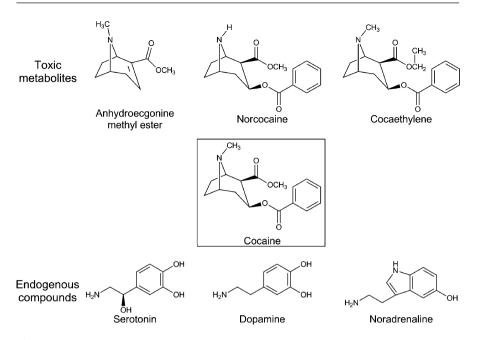
In the streets, cocaine may be available in water-soluble or insoluble forms. Watersoluble forms include cocaine sulfate and cocaine hydrochloride, which are mainly used intranasally, but may also be administered orally, sublingually, or by injection. Some drug abusers inject cocaine hydrochloride subcutaneously, intramuscularly, or intravenously, alone or with heroin ("speedball") or other drugs. Water-insoluble forms such as free base cocaine or crack are usually smoked. Crack cocaine is abused by inhaling the vapor from cigarettes (usually mixed with tobacco or marijuana) or after heating the drug in a glass pipe. When the drug is consumed in the free base form, thermal degradation leads to the formation of the metabolite anhydroecgonine methyl ester (Fig. 1), recently shown to be neurotoxic (Garcia et al. 2012).

Upon consumption, cocaine rapidly reaches the brain because it is able to cross the blood–brain barrier, due to its high lipophilicity, in contrast to its main metabolites ecgonine methyl ester and benzoylecgonine (Buttner et al. 2003). However, cocaine may be converted into other toxic metabolites, such as norcocaine and cocaethylene (Fig. 1). Norcocaine is formed in the liver, via N-demethylation of cocaine by CYP3A4, and is an active cocaine metabolite that crosses the blood–brain barrier and induces neurotoxicity (Nassogne et al. 1997). In the presence of alcohol, another active metabolite, cocaethylene (Fig. 1), is formed, which is more toxic than cocaine itself (Henry 2007), as discussed in Sect. 6.2.

Cocaine available in the streets is often adulterated with sugars, other central nervous system stimulants and local anesthetics (Karch 2009), to increase profit or to modulate the effects experienced by its users, being an additional factor that may contribute to some of the neuropathology observed in cocaine abusers. Purity of cocaine generally ranges from 20 % to 95 %, for cocaine hydrochloride, and 20 % to 80 %, for crack (Couper and Logan 2004).

### 3 Neurologic Dysfunction in Cocaine Abusers

Cocaine addiction may be considered a neurological disorder (Majewska 1996) because several neurological impairments are found in cocaine abusers, including seizures, cerebral ischemia, cerebral hemorrhages, infarction, optic neuropathy, cerebral atrophy, cognitive impairment, and mood and movement disorders (Koppel et al. 1996; Majewska 1996). Movement disorders described in cocaine users include choreoathetosis, akathisia, and parkinsonism with tremor (Karch 2009), a phenomenon known as "crack dancing," which may be associated with



**Fig. 1** Chemical structures of cocaine, its toxic metabolites, and monoamine endogenous compounds that share structural similarities with the drug. These similarities are in the basis of many of the drug's effects, due to interference with the same endogenous molecular targets

an increased incidence of basal ganglia abnormalities in cocaine abusers (Bartzokis et al. 1999).

The prefrontal cortex and basal ganglia of cocaine abusers are the brain regions where most brain lesions and cerebral atrophy are observed (Bartzokis et al. 1996; Langendorf et al. 1996). Cocaine abusers also present specific dysfunction of executive functions associated with prefrontal brain regions (Bolla et al. 1998). Cocaine-induced brain dysfunction may be caused by (i) vasoconstriction and consequent hypoxia induced by the drug (Olsen 1995); (ii) impairment of neuro-transmission, namely through dopamine and glutamate; and (iii) oxidative stress and mitochondrial dysfunction.

The neurotoxic actions of cocaine in different brain areas involve different mechanisms of action and have specific consequences. Cocaine-induced increase in dopamine activity in the limbic system translates into euphoria, confusion, agitation, and hallucination. Effects in the cortex lead to pressure of speech, excitation, and a reduced feeling of fatigue. Stimulation of lower centers leads to tremor and tonic-clonic convulsions. Cocaine effects in the brain stem lead to stimulation and then depression of the respiratory vasomotor and vomiting centers (International Programme on Chemical Safety 1999). Maternal abuse of cocaine is associated with prenatal brain toxicity, leading to structural, metabolic, and functional brain abnormalities in the offspring (Roussotte et al. 2010).

### 4 Impairment of Neurotransmission

As most drugs of abuse, cocaine affects neurotransmission in brain areas involved in reward, such as the ventral pallidum; in motivation/drive, such as the orbitofrontal cortex and the subcallosal cortex; in memory and learning, such as the amygdala and the hippocampus; and in impulse control, such as the prefrontal cortex and the anterior cingulate gyrus (Volkow et al. 2003). The addictive effects of cocaine, and virtually all drugs of abuse, are explained by the activation of the brain reward pathway, a neuronal circuit composed of dopaminergic neurons that project from the ventral tegmental area, in the midbrain; to the nucleus accumbens, in the ventral striatum; and to the prefrontal cortex (Hyman et al. 2006).

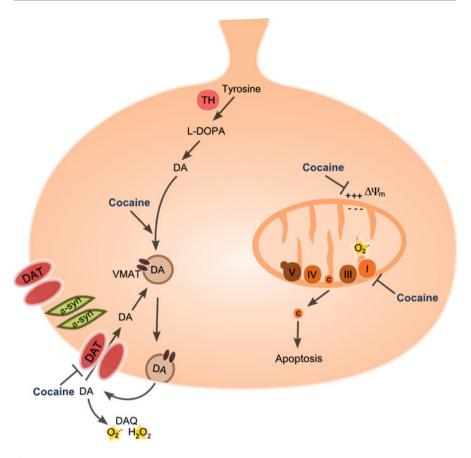
# 4.1 Cocaine Effects on Dopamine and Glutamate Signaling

Due to structural similarities with monoamines (Fig. 1), cocaine interferes with the reuptake of these neurotransmitters, particularly dopamine, which is mostly associated with pleasure and movement. Blockade of monoamine reuptake (Ki: 02–0.7  $\mu$ M) explains most of the effects of cocaine on the central nervous system (Han and Gu 2006). Additionally, at higher concentrations (>5  $\mu$ M), cocaine may also bind to voltage-gated ion channels and inhibit inward sodium and outward potassium currents, acting as a local anesthetic (Heard et al. 2008). These effects are relevant for the range of cocaine concentrations found in the plasma of cocaine addicts, usually between 0.3 mM and 1 mM, which may even represent higher cocaine concentrations in the brain, because brain cocaine concentrations were reported to exceed the plasma concentration by several times (Heard et al. 2008).

By binding to the dopamine transporter (DAT) (Fig. 2), cocaine blocks its function, leading to increased extracellular dopamine, which results in chronic stimulation of postsynaptic dopamine receptors (Cunha-Oliveira et al. 2008, for review) and leads to the euphoric "rush." Subsequently, dopamine levels fall, resulting in the dysphoric "crash." Cocaine also interferes with the uptake of norepinephrine and serotonin, leading to increased stimulation of postsynaptic receptors for these neurotransmitters (Heard et al. 2008).

Besides inhibiting plasma membrane monoamine transporters, cocaine also interacts with the vesicular monoamine transporter VMAT-2 (Fig. 2), favoring the storage of catecholamines inside synaptic vesicles (Brown et al. 2001). Concomitant inhibition of the DAT and increased vesicular sequestration of dopamine induced by cocaine may cause a shift in the ratio of cytosolic to vesicular dopamine, increasing the amount of neurotransmitter packaged in each vesicle before its release, which may contribute to a greater increase in synaptic dopamine accumulation upon a depolarizing stimulus (Brown et al. 2001).

Several brain structures that receive input from the reward pathway send reciprocal glutamatergic projections back to the ventral tegmental area, which can affect dopamine release (Uys and Reissner 2011). Glutamatergic neurotransmission has been implicated in addiction, in the processes of reinforcement, sensitization, habit



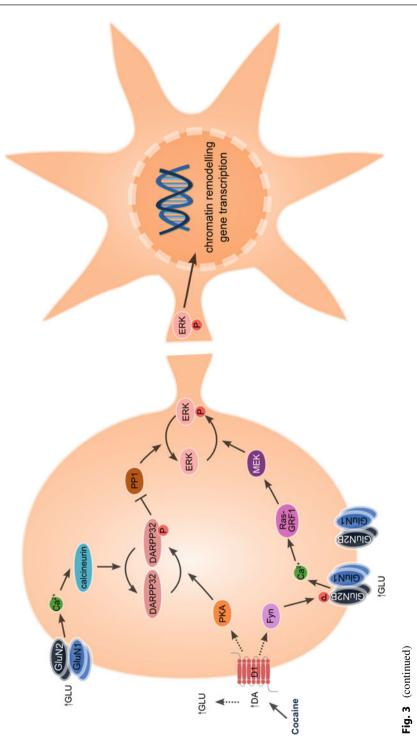
**Fig. 2** Presynaptic changes induced by acute cocaine exposure. Cocaine affects neurotransmission in dopaminergic and other monoaminergic nerve terminals. Cocaine inhibits the dopamine (DA) transporter (DAT), preventing the reuptake of DA into the nerve terminal and prolonging the effects of this neurotransmitter in the synapse. Cocaine also interacts with the vesicular monoamine transporter (VMAT)-2, favoring the storage of catecholamines inside synaptic vesicles, which may contribute to a greater increase in synaptic dopamine, upon a depolarizing stimulus. DA is easily auto-oxidized, generating DA quinone (DAQ) and reactive oxygen species, such as  $O_2^{--}$  and  $H_2O_2$ . Cocaine exposure also affects alpha-synuclein and DAT expression. These two proteins may interact, modulating DA neurotransmission. In addition, since cocaine is positively charged at physiological pH, it may enter into the nerve terminal and interact directly with mitochondria and other intracellular targets, inhibiting complex I-driven mitochondrial respiration, dissipating mitochondrial potential, and activating the mitochondrial apoptotic pathway

learning, context conditioning, craving, and relapse (Tzschentke and Schmidt 2003); moreover, glutamate receptors were shown to play a role in the reinforcement of long-lasting drug-seeking behaviors (Uys and Reissner 2011). Glutamatergic effects of drugs of abuse in the reward pathway may be modulated by dopamine in the nucleus accumbens, which controls the efficacy of glutamatergic corticostriatal synapses (Wickens et al. 2007). Chronic cocaine exposure induces synaptic plasticity at ventral tegmental area and nucleus accumbens glutamatergic synapses, including changes in structural plasticity (i.e., increase in the number of dendritic spines); in glutamate homeostasis, implicating glial and neuronal impairment; and in postsynaptic glutamate signaling (Uys and Reissner 2011).

Cocaine was shown to increase extracellular glutamate concentrations in brain areas such as the ventral tegmental area, nucleus accumbens, prefrontal cortex, or striatum (Williams and Steketee 2004). An increase in extracellular glutamate has been associated with excitotoxic processes, due to increased activation of *N*-methyl-D-aspartate (NMDA) receptors and subsequent increase in intracellular  $Ca^{2+}$  concentration ( $Ca^{2+}_{i}$ ) (Rego and Oliveira 2003, for review), which could contribute to cocaine neurotoxicity. Indeed, an increase in  $Ca^{2+}_{i}$  was observed in rat cortical neurons, after cocaine exposure (Cunha-Oliveira et al. 2010).

Plasticity of glutamatergic synaptic transmission in the ventral tegmental area contributes to the development of addictive behaviors in cocaine addicts (Schilstrom et al. 2006; Wolf 2010). Changes in NMDA receptor subunit expression were observed in the brains of cocaine-exposed rats (Huber et al. 2001; Schilstrom et al. 2006; Yamaguchi et al. 2002; Scheggi et al. 2002; Hemby et al. 1999). In neurons from the ventral tegmental area, cocaine was shown to increase the expression of GluN1 and GluN2B (which is mainly extrasynaptic) subunits and their redistribution to synaptic membranes (Schilstrom et al. 2006). GluN2B expression was also shown to increase in the nucleus accumbens and hippocampus of cocaine-exposed rats. This effect was prevented in rats exposed to cocaine and MK-801, an NMDA receptor antagonist (Scheggi et al. 2002). Altered composition of NMDA receptors may increase the susceptibility of cells to the toxicity of drugs of abuse, as demonstrated by the fact that GluN1/GluN2B-transfected HEK293 cells are more susceptible to acute street heroin toxicity when compared to GluN1/GluN2A- and GluN1-transfected cells (Domingues et al. 2006).

The changes in the NMDA receptor subunit expression induced by cocaine may be mediated by extracellular dopamine, via stimulation of dopamine receptors (Schilstrom et al. 2006). Indeed, dopamine receptors D2 and D1 were shown to interact with NMDA receptor subunits, contributing to the stimulant effect of cocaine. Interactions between GluN2B subunits of the NMDA receptor and the D2 dopamine receptor were observed in the neostriatum of cocaine-exposed rats (Liu et al. 2006). In addition, D1 dopamine receptors, NMDA receptors, and extracellular-signal-regulated kinase (ERK) significantly contributed to neuronal morphological changes induced by repeated exposure to cocaine (Fig. 3) (Ren et al. 2010). Activation of NMDA receptors in the striatum was shown to recruit cytoplasmic D1 receptors to the plasma membrane and dendritic spines (Missale et al. 2006; Sun et al. 2008). D1 and NMDA receptors regulate, in opposite directions, the extent of dopamine- and cAMP-regulated neuronal phosphoprotein (DARPP32) phosphorylation, which becomes a potent inhibitor of protein phosphatase 1 (PP1) (Fig. 3). Whereas activation of the D1 receptor leads to increase in cAMP formation and activation of protein kinase A (PKA), which phosphorylates DARPP32, the stimulation of NMDA receptors induces an increase in intracellular calcium, leading



to the activation of calcineurin and dephosphorylation of DARPP32 (Nakano et al. 2010). Interestingly, interaction between glutamate and dopamine signaling in striatal medium spiny neurons seems to be critical for long-term plasticity in the striatum and may play a role in the behavioral alterations induced by cocaine (Pascoli et al. 2011). These adaptations seem to depend on activation of the ERK pathway in the striatum (Ren et al. 2010), which has been pointed out as an essential event for chromatin remodeling and gene expression associated with drug abuse (Brami-Cherrier et al. 2009). Interestingly, ERK activation upon cocaine exposure occurs selectively in dopamine D1 receptor-expressing medium spiny neurons and requires the coincident stimulation of D1 and NMDA receptors (Pascoli et al. 2011). D1 and NMDA receptor interplay upon cocaine exposure may result in ERK activation in medium spiny neurons and may be mediated by two pathways (Fig. 3). Stimulation of D1 receptors leads to PKA activation and DARPP-32 phosphorylation, inhibiting PP1 and preventing ERK dephosphorylation. Alternatively, D1 receptor activation also mediates the phosphorylation and activation of the Src-family kinase Fyn, which phosphorylates GluN2B subunits of NMDA receptors, increasing the entry of Ca<sup>2+</sup> through these receptors, which leads to the activation of Ras protein-specific guanine nucleotide-releasing factor 1 (Ras-GRF1), mitogen-activated protein kinase kinase (MAPKK)/MEK, and ERK, representing a powerful mechanism of NMDA receptor potentiation (Pascoli et al. 2011).

# 5 Mechanisms of Neurotoxicity

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Cocaine neuropathology shares some common aspects with neurodegenerative disorders, such as Parkinson's disease (PD), which also involves a dopaminergic dysfunction. As described in Sect. 3, chronic cocaine use leads to movement disorders, with similarities to parkinsonism (Karch 2009), although cocaine use does not seem to be a risk factor for the development of idiopathic PD (San Luciano and Saunders-Pullman 2009; Callaghan et al. 2012).

**Fig. 3** Postsynaptic signaling in glutamate/dopamine-responsive striatal medium spiny neurons, upon chronic cocaine exposure. By increasing synaptic dopamine (*DA*) and glutamate (*GLU*), cocaine induces changes in the expression of NMDA receptor subunits and alterations in post-synaptic signaling through DARPP-32 and ERK. Stimulation of the D1 receptor leads to PKA activation, which phosphorylates DARPP-32, with subsequent inhibition of PP1, thus preventing ERK dephosphorylation. NMDA receptors also modulate DARPP-32 phosphorylation, since intracellular calcium entering through the receptor activates the phosphatase calcineurin, which dephosphorylates DARPP-32, preventing PP1 inhibition and enabling subsequent ERK dephosphorylation. Cocaine-induced ERK activation occurs selectively in D1 receptor-expressing MSNs and requires the coincident stimulation of D1 and NMDA receptors. This may be explained by the activation of Fyn following D1 receptor stimulation, which leads to phosphorylation of GluN2B subunits, resulting in increased calcium permeability, leading to the activation of D1 receptors may lead to ERK activation by two independent pathways. Active ERK mediates chromatin remodeling and gene transcription that may be involved in behavioral manifestations of cocaine addiction

Cocaine exposure affects the expression of alpha-synuclein, the first protein found to be mutated in genetic cases of PD, whose accumulation in intracellular filamentous aggregates constitutes a pathological feature of PD and other neurodegenerative disorders (Perfeito et al. 2012). An overexpression of alpha-synuclein was found in dopamine neurons of cocaine abusers (Mash et al. 2003) and in the nucleus accumbens of rats exposed to high-dose cocaine (Brenz Verca et al. 2003). Increased alpha-synuclein protein levels were also found in serum from recently abstinent cocaine abusers (Mash et al. 2008). Interestingly, alpha-synuclein was reported to interact with the DAT, a cocaine classic target, modulating dopamine transmission and neuronal function (Swant et al. 2011; Bellucci et al. 2011). The presence of elevated levels of alpha-synuclein and DAT in the striatum of human cocaine abusers (Qin et al. 2005) suggested that cocaine may concomitantly regulate the binding and function of these two proteins (Fig. 3). Thus, overexpression of alpha-synuclein induced by cocaine may play a role in cocaineinduced plasticity and regulation of dopamine synaptic tone (Qin et al. 2005). Accordingly, alpha-synuclein overexpression in rat nucleus accumbens was suggested to modulate cocaine-induced locomotion and self-administration (Boyer and Dreyer 2007).

Other common characteristics observed in cocaine abuse and in several neurodegenerative diseases include oxidative stress, mitochondrial dysfunction, and apoptosis, as described in the next sections.

# 5.1 Cocaine Toxicity and Oxidative Stress

The occurrence of oxidative stress in neurons upon cocaine exposure has been suggested to occur due to the presence of oxidized metabolites of cocaine (Kovacic 2005), due to an accumulation of dopamine and its metabolites in the brain, or due to the deregulation of antioxidant defenses (Cunha-Oliveira et al. 2008, for review), all of which may ultimately result in increased levels of reactive oxygen species.

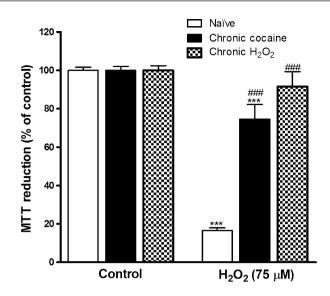
As described in Sect. 4.1, cocaine blocks dopamine reuptake by the DAT, resulting in the increase in extracellular dopamine levels in specific brain areas. Dopamine has been shown to be neurotoxic in vitro (Graham et al. 1978; McLaughlin et al. 1998) and in vivo (Hastings et al. 1996). Dopamine has a propensity to form reactive metabolites by enzymatic and nonenzymatic mechanisms and may induce oxidative stress in dopaminergic and neighboring cells, which may contribute to the neurotoxicity of many drugs of abuse, particularly cocaine due to its direct effect in increasing synaptic dopamine oxidase A (MAO<sub>A</sub>) (Fornai et al. 2000) and in a lower extent by MAO<sub>B</sub> (Youdim et al. 2006), two isoforms of a mitochondrial enzyme that is present in the cytoplasmic side of the outer mitochondrial membrane in neurons and astrocytes. This enzyme generates 3,4-dihydroxyphenylacetaldehyde (DOPAL), which is highly toxic and rapidly metabolized by aldehyde dehydrogenase, producing 3,4-dihydroxyphenylacetic acid (DOPAC) and hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) (Marchitti et al. 2007). In contrast

with enzymatic dopamine oxidation, dopamine auto-oxidation may also occur extracellularly, which is more relevant for explaining cocaine-induced oxidative stress. Auto-oxidation of the catechol ring of dopamine generates dopamine quinones, together with superoxide anion  $(O_2^{\bullet-})$  and  $H_2O_2$  (Fig. 2), which may react with transition metal ions, such as iron, via the Haber-Weiss/Fenton reactions, creating the highly toxic hydroxyl radical ( $^{\bullet}OH$ ) (Hastings 2009).  $O_2^{\bullet-}$  may also lead to the formation of the highly toxic peroxynitrite, by reaction with nitric oxide. Dopamine quinones are electron-deficient molecules and, thus, readily react with cellular nucleophiles, such as reduced sulfhydryl groups on protein cysteinyl residues, covalently modifying protein structure. Since cysteinyl residues are often localized at the active site of proteins, covalent modification by dopamine quinones often leads to inactivation of protein function, affecting cell survival (Hastings 2009).

Cocaine also affects antioxidant defenses, which protect the cells against free radicals and other reactive species, such as  $O_2^{\bullet-}$  and  $H_2O_2$ . Superoxide dismutase (SOD) is an antioxidant enzyme that detoxifies  $O_2^{\bullet-}$ , but may contribute to increase  $H_2O_2$  levels (Cunha-Oliveira et al. 2008, for review). The main antioxidant enzymes involved in  $H_2O_2$  inactivation are glutathione peroxidase (GPx) and catalase, which is present in peroxisomes. These enzymes are regulated by the cells, through the control of protein expression or activity, which allows the maintenance of cellular homeostasis and leads to a contrast in the cellular effects of chronic and acute oxidant exposures. For example, whereas acute exposure to  $H_2O_2$  induces apoptotic cell death (Benedi et al. 2004; Jang and Surh 2004), chronic exposure to low  $H_2O_2$  concentrations induces cellular resistance to the acute toxicity of this compound (Cunha-Oliveira et al. 2006b).

Acute cocaine exposure increased  $H_2O_2$  in the prefrontal cortex and in the striatum of rats (Dietrich et al. 2005) and decreased catalase activity in the same structures in mice (Macedo et al. 2005). The levels of the antioxidants vitamin E and reduced glutathione (GSH) were also shown to be decreased in human neuronal progenitor cells upon acute cocaine exposure (Poon et al. 2007) and in fetal rat brain upon in utero cocaine exposure (Lipton et al. 2003). Biochemical markers of oxidative injury were also found in brain cells, after cocaine exposure. The hippocampus of rats exposed prenatally to cocaine showed an increase in lipid peroxidation (Bashkatova et al. 2006), and protein oxidation was found in human neuronal progenitor cells acutely exposed to cocaine (Poon et al. 2007). Elevated malondialdehyde (MDA) and nitrite levels were also found in rat brain slices from the prefrontal cortex and nucleus accumbens, after acute cocaine exposure. This increase in oxidative damage markers was accompanied by a decrease in total antioxidant content, and both were prevented by tempol, a SOD mimetic antioxidant (Numa et al. 2008).

Repeated cocaine exposure decreased GSH concentration and GPx activity in the hippocampus of rats, which also presented learning and memory impairments (Muriach et al. 2010). Interestingly, chronic exposure of catecholaminergic cells to cocaine seems to induce adaptation to oxidative stress, because PC12 cells chronically exposed to cocaine exhibited a partial resistance against  $H_2O_2$  toxicity (Fig. 4) (Cunha-Oliveira et al. 2006b), which suggests the occurrence of oxidative stress

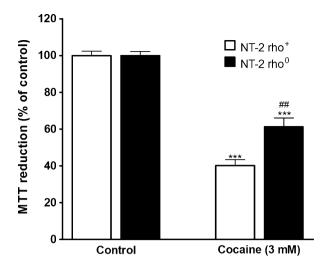


**Fig. 4** Involvement of oxidative stress in chronic cocaine effects. PC12 cells were chronically exposed to subtoxic concentrations of cocaine (30  $\mu$ M) or H<sub>2</sub>O<sub>2</sub> (10  $\mu$ M) for 7–12 months. The toxicity of an acute exposure to H<sub>2</sub>O<sub>2</sub> was then assessed by the MTT assay. Cells chronically exposed to H<sub>2</sub>O<sub>2</sub> were completely resistant to H<sub>2</sub>O<sub>2</sub> acute toxicity, due to cellular adaptation to oxidative stress. Cells chronically exposed to cocaine were partially resistant to H<sub>2</sub>O<sub>2</sub> toxicity, suggesting that chronic cocaine exposure also induces cellular adaptation to oxidative stress. Statistical analysis was performed by one-way ANOVA with Bonferroni posttest \*\*\**P* > 0.001 compared to respective control; ###*P* > 0.001 compared to treated naïve cells (Adapted from Toxicology, 217, T. Cunha-Oliveira, A.C. Rego, M.T. Morgadinho, T. Macedo and C.R. Oliveira, Differential cytotoxic responses of PC12 cells chronically exposed to psychostimulants or hydrogen peroxide, 54–62 (Copyright 2006. Cunha-Oliveira et al. (2006b), with permission from Elsevier)

during chronic cocaine exposure. In dopaminergic rat brain structures, repeated cocaine administration induced an increase in antioxidant enzyme activity (Dietrich et al. 2005), and repeated self-administration of cocaine in rats induced an increase in SOD activity and a reduction in MDA levels in the hippocampus, frontal cortex and dorsal striatum (Pomierny-Chamiolo et al. 2012). Induction of antioxidant defenses in models of repeated cocaine exposure may explain the resistance to oxidative agents in these models, and cocaine-induced adaptations in cellular redox balance were suggested to contribute to enduring behavioral plasticity (Uys et al. 2011).

#### 5.2 Cocaine Toxicity and Mitochondrial Dysfunction

Mitochondrial function and energy metabolism were shown to be affected in brains of human cocaine abusers (Lehrmann et al. 2003), and cocaine was demonstrated to influence mitochondrial function in different cell types. Interestingly, acute cocaine



**Fig. 5** Partial involvement of functional mitochondria in cocaine toxicity. Mitochondrial involvement in acute cocaine toxicity was analyzed in NT-2 rho-zero cells, depleted from mitochondrial DNA, and thus devoid of a functional mitochondrial respiratory chain. Loss of cell viability induced by cocaine was attenuated in NT-2 rho-zero cells, compared with NT-2 rho-plus cells. Statistical analysis was performed by one-way ANOVA with Bonferroni posttest \*\*\**P* > 0.001 compared to respective control; ##*P* > 0.01 compared to treated NT-2 rho-plus cells (Adapted from Brain Research, 1089, T. Cunha-Oliveira, A.C. Rego, S.M. Cardoso, F. Borges, R.H. Swerdlow, T. Macedo and C.R. Oliveira, Mitochondrial dysfunction and caspase activation in rat cortical neurons treated with cocaine or amphetamine, 44–54. Copyright 2006. Cunha-Oliveira et al. (2006a), with permission from Elsevier)

toxicity partially required the presence of a functional respiratory chain, because rho-zero cells, which lack functional mitochondria, were more resistant to cocaine toxicity, in comparison with rho-plus cells (Fig. 5) (Cunha-Oliveira et al. 2006a). In addition, exposure to cocaine downregulated mitochondrial gene expression in rat cingulate cortex (Dietrich et al. 2004). Accordingly, mitochondrial complex I activity and subunit expression were reduced after cocaine exposure (Yuan and Acosta, Jr. 2000; Dietrich et al. 2004; Devi and Chan 1997), a defect that is also observed in PD. Cocaine may interact directly with mitochondria and other intracellular targets (Heard et al. 2008), entering the cell due to the weak base effect, because cocaine has a positive charge at physiological pH. Studies in hepatic mitochondria showed that in vivo cocaine administration decreased state 3 respiration, the respiratory control ratio, and the activity of complexes I, II/III, and IV (Devi and Chan 1997). In isolated liver and brain mitochondria, cocaine inhibited complex-I-driven respiration, through a direct effect on this complex (Cunha-Oliveira et al. 2013).

Cocaine neurotoxicity has been associated with apoptotic cell death, and cocaine has been shown to induce biochemical features of apoptosis, including activation of caspases, loss of mitochondrial potential, and cytochrome c release (Cunha-Oliveira et al. 2008, for review). In brains of cocaine addicts, the occurrence of

aberrant cell death was suggested by the enhanced degradation of nuclear poly (ADP-ribose) polymerase (PARP)-1, an apoptotic hallmark, which appeared to be the consequence of oxidative stress and activation of nuclear AIF (Alvaro-Bartolome et al. 2011). Changes in Bcl-2 and Bax expression (Lepsch et al. 2009; Dey et al. 2007), decrease of mitochondrial cytochrome c levels, and caspase-9 activation (Cunha-Oliveira et al. 2006a) were found after in vitro cocaine exposure, suggesting the involvement of the mitochondrial apoptotic pathway. However, controversial results have been reported regarding structural apoptotic hallmarks induced by cocaine. Whereas many reports did not find structural features of neuronal apoptosis upon in vivo (Dietrich et al. 2005) or in vitro (Oliveira et al. 2006a) cocaine exposure, others have shown that cultured fetal mouse cortical neurons exposed to cocaine exhibited apoptotic morphology (Nassogne et al. 1997) and apoptotic neurons were also observed in the fetal rat brain, upon in utero cocaine exposure, whereas the maternal brain was spared (Xiao and Zhang 2008).

Interestingly, oxidative stress seemed to precede cell death induced by cocaine, in human neuronal progenitor cells (Poon et al. 2007). Oxidative stress also seems to be involved in cocaine's mitochondrial effects, since cocaine-induced mitochondrial dysfunction could be prevented by the mitochondrial-targeted antioxidant MitoQ, in isolated cardiac mitochondria (Vergeade et al. 2010).

# 6 Combinations of Cocaine and Other Substances

# 6.1 Effects of Speedball

Cocaine abusers may also consume other drugs that may interact with cocaine and modify its effects. Speedball is a relatively popular drug combination (European Monitoring Center for Drugs and Drug Addiction 2009), which consists in simultaneous injection of cocaine and heroin, and has been reported to cause more rewarding effects in rats than cocaine or heroin alone (Ranaldi and Munn 1998). The popularity of this drug combination may derive from the reduction of the unwanted side effects of one drug by the other due to compensatory mechanisms of action and/or from enhanced effects of the combination (Leri et al. 2003).

Speedball has serious health consequences. The use of cocaine in combination with heroin is generally associated with the presence of a mental illness and may aggravate underlying psychological problems, such as bipolar disorder (European Monitoring Center for Drugs and Drug Addiction 2008). Moreover, speedball abusers seem to exhibit more severe psychopathology in comparison with other cocaine addicts and more likely fail in drug abuse treatment (Bandettini Di Poggio et al. 2006).

Self-administration of cocaine in rhesus monkeys was shown to be enhanced by otherwise inactive doses of heroin (Rowlett and Woolverton 1997). The pharma-cological mechanisms underlying speedball abuse seem to involve an increase in the reward obtained from the drug combination (Rowlett and Woolverton 1997).

In accordance, self-administration of both cocaine and heroin produces synergistic elevations in extracellular dopamine concentration in the reward pathway (Hemby et al. 1999; Smith et al. 2006) that are thought to be mediated by dopamine and  $\mu$ -opioid receptors (Cornish et al. 2005). This suggests that this drug combination alters the dynamics of dopamine and other neurotransmitters affected in drug addiction.

Interestingly, a chemical interaction between cocaine and morphine was found in cocaine-heroin mixtures, involving the formation of a cocaine-morphine adduct (Garrido et al. 2007), which may play a role in the effects of cocaineheroin combinations, especially when the co-abuse occurs simultaneously. Thus, besides pharmacodynamic interactions, chemical interactions between drugs could also play a role in speedball neurotoxicity. In previous studies, we exposed primary cultures of rat cortical neurons to heroin and/or cocaine, either sequentially or simultaneously (Cunha-Oliveira et al. 2010) to investigate this possibility. The effects of cocaine seemed to predominate over heroin's effects when the cells were exposed to a mixture of the two drugs (where the probability of chemical interactions between the drugs is enhanced), but not upon sequential exposure (Cunha-Oliveira et al. 2010). Since both drugs were present at the same amounts in both situations, this result suggested that the interaction of the mixture of drugs with opioid receptors may be lower, as compared to a sequential administration of the two drugs, probably due to the presence of cocaine-morphine adducts. In addition, cocaine-morphine combinations induced apoptosis and mitochondrial dysfunction in rat cortical neurons (Cunha-Oliveira et al. 2010) and impaired the bioenergetics of isolated brain mitochondria (Cunha-Oliveira et al. 2013). These data suggest that neurotoxic pathways induced by coadministration of cocaine and heroin differ from those evoked by a sequential administration of the same drugs, possibly due to chemical interactions between the two drugs (or their metabolites), which may interfere with cell death mechanisms and may affect the capacity of cocaine to interact with its cellular targets, such as monoamine transporters or mitochondria. Thus, co-use of cocaine and heroin may lead to different effects depending on the mode of co-exposure, and polydrug abusers may be more prone to neurotoxic damage than single drug abusers.

# 6.2 Cocaine and Ethanol

As referred in Sect. 2, cocaine and ethanol represent another toxic drug combination. Ethanol use by cocaine abusers leads to the potentiation of cocaine's euphoric effects. However, toxic effects of both drugs are also enhanced (Bolla et al. 2000), since the drugs are combined in the liver to form a very toxic metabolite – cocaethylene (see Fig. 1). Cocaethylene also affects the brain because it is highly lipophilic and, thus, crosses the blood–brain barrier. The behavioral effects of cocaethylene are similar to those of cocaine, but more prolonged due to its longer half-life (Henry 2007).

### 7 Conclusion

Cocaine is a widely abused psychostimulant drug, popularized by its intense euphoric effects. Cocaine addiction may be described as a neurological disorder because cocaine abusers present several neurological impairments. Cocaine neurotoxicity may be due to vasoconstriction and subsequent hypoxia or to the impairment of neurotransmission, namely through dopamine and glutamate, and may be mediated by oxidative stress and mitochondrial dysfunction. The interaction with the dopaminergic system is the main cause of cocaine's addictive effects, leading to excess synaptic dopamine levels that may induce oxidative stress through dopamine auto-oxidation and reactive oxygen species generation. Neurotoxic cocaine metabolites may also induce oxidative stress. Cocaine also affects glutamate levels and the expression of glutamate receptors, which may mediate excitotoxic cell damage and long-term neuroadaptive effects.

Cocaine abuse shares some common aspects with neurological disorders such as PD, which also involves a dysfunction in the dopaminergic system. Cocaine exposure has been shown to regulate both alpha-synuclein and DAT binding and function, possibly affecting cocaine-induced plasticity, locomotion, and self-administration. Moreover, cocaine has been shown to impair mitochondrial function in several models, namely through the inhibition of mitochondrial respiratory chain complex I, as occurs in PD; furthermore, cocaine toxicity was demonstrated to require a functional respiratory chain. Cocaine may also induce the activation of mitochondrial apoptotic pathways in several models.

Cocaine is frequently co-abused with heroin, in a combination known as speedball, and heroin may modulate or modify cocaine's effects. A chemical interaction between cocaine and morphine was found in drug mixtures similar to those used by speedball abusers and may produce a specific neurotoxic profile and alter cocaine effects. A concomitant use of cocaine and alcohol may have additive negative effects in the brain, due to the formation of a metabolite, cocaethylene, which is more toxic than cocaine itself.

In summary, cocaine-induced neurotoxicity may underlie brain dysfunction in cocaine and polydrug abusers and may predispose the brain to neurodegeneration.

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# 5,6- and 5,7-Dihydroxytryptamines as Serotoninergic Neurotoxins

# Justyna Paterak and Roman Stefański

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e-mail: jpaterak@ipin.edu.pl; stefans@ipin.edu.pl

J. Paterak • R. Stefański (🖂)

Department of Pharmacology and Physiology of the Nervous System, Institute of Psychiatry and Neurology, Warsaw, Poland

### Abstract

Dihydroxytryptamines are able to selectively destroy central serotonin neurons when catecholaminergic neurons are protected. Compared to serotonin, 5,6dihydroxytryptamine (5,6-DHT) and 5,7-dihydroxytryptamine (5,7-DHT) are indole derivatives that acquire their neurotoxicity due to additional hydroxyl groups on the indole ring. Factors influencing the action of dihydroxytryptamines include the site of injection, the speed and volume of injection, the amount and type of antioxidant used, the type of anesthesia, and the dose and type of catecholamine uptake blocker. Two major hypotheses have been proposed to explain the molecular mechanism that underlies the toxicity of 5,6- and 5,7-DHT. The first suggests that highly reactive (electrophilic) quinonoid intermediates, created as a result of 5,6- and 5,7-DHT auto-oxidation, bind covalently to molecules indispensable for neuronal function. The second hypothesis states that during the auto-oxidation of dihydroxylated tryptamines, reactive oxygen species such as superoxide radical anion  $(O_2^{-})$ , hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), and the hydroxyl radical (HO) are generated as toxic byproducts. The use of 5,6- and 5,7-DHT in both adult and neonatal animals enables the recognition the role of 5-HT in locomotor activity (LMA), prepulse inhibition, seizure susceptibility, sleeping behavior, feeding and drinking behavior, sexual and aggressive behavior, the regulation of body temperature, the response to novel and noxious stimuli, learning and memory, the antinociceptive action of drugs, neuroendocrine regulatory mechanisms, and developmental plasticity.

#### Keywords

5,6-dihydroxytryptamine (5,6-DHT) • 5,7-dihydroxytryptamine (5,7-DHT) • 5-HT neurotoxins

List of Abbreviations						
5-HT	5-hydroxytryptamine, serotonin					
5-HTP	5-hydroxytryptophan					
5,6-DHT	5,6-dihydroxytryptamine					
5,7-DHT	5,7-dihydroxytryptamine					
6-OHDA	6-hydroxydopamine					
CNS	Central nervous system					
DA	Dopamine					
DRN	Dorsal raphe nuclei					
DTPA	Diethylene triamine pentaacetic acid					
FSH	Follicle-stimulating hormone					
GH	Growth hormone					
GSH	Glutathione					
$H_2O_2$	Hydrogen peroxide					
HO	Hydroxyl radical					

L-5-HTP	5-hydroxy-L-tryptophan
LMA	Locomotor activity
MAO	Monoamine oxidase
MAOI	Monoamine oxidase inhibitor
MRN	Median raphe nuclei
NA	Noradrenaline
NGF	Nerve growth factor
$0_{2}^{-}$	Superoxide radical anion
PPI	Prepulse inhibition
PTZ	Pentylenetetrazol
REM	Rapid eye movements
ROS	Reactive oxygen species
SOD	Superoxide dismutase

### 1 Introduction

# 1.1 The Significance of Serotonin and Serotonergic Neurons

Serotonergic neurons are one of the first to appear during brain and spinal cord development. They are formed very early in gestation in vertebrates (Lidov and Molliver 1982). Serotonin (5-hydroxytryptamine or 5-HT) is a neurotransmitter with developmental functions (Ahmad and Zamenhof 1978) and is involved in more behaviors, physiological processes, and diseases than any other transmitter. It affects developmental events such as cell division, neuronal migration, cell differentiation, and synaptogenesis (Lipton and Kater 1989; Azmitia 2001; Vitalis and Parnavelas 2003). Studies in vertebrates show that even a brief 5-HT deprivation in the brain leads to a loss of spines, dendritic profiles, and synapses (Yan et al. 1997).

# 1.2 From the First Neurotoxins to the Discovery of Dihydroxytryptamines

The roles of different neuronal types and pathways can be determined by disrupting their function. In the first half of the twentieth century, using the methods of axonal transaction and chemical or electrolytic ablation of nuclei, our knowledge of nerve function was enhanced. However, these techniques have some inherent limitations due to the lack of selectivity. A real break-through came in the 1950s with the discovery the first neurotoxin, an antibody to nerve growth factor (NGF), which could destroy sympathetic nerves in

whole animals. The next neurotoxin introduced in the 1960s by Thoenen and Tranzer (1968) was 6-hydroxydopamine (6-OHDA). This hydroxylated analog of dopamine (DA) promoted selective degeneration of peripheral and central noradrenergic neurons. This discovery initiated a novel period in experimental biology. Selective chemical axotomy replaced early mechanical and electrolytic lesioning techniques. After the discovery of 6-OHDA, Baumgarten and collaborators (1971) concentrated on dihydroxytryptamine derivatives. They found that these compounds are able to cause selective destruction of central serotonergic neurons when noradrenergic neurons are protected through pharmacological inhibition of the noradrenaline reuptake carrier (Baumgarten and Lachenmayer 1972; Baumgarten et al. 1973). In 1978, Jacoby and Lytle first used the term "serotonin neurotoxins" in the title of their symposium proceedings volume to describe these compounds.

The mechanism of action of dihydroxytryptamines is analogous to the effects triggered by 6-OHDA in catecholaminergic terminals. Initially, 5,6-dihydroxytryptamine (5,6-DHT) was introduced to neuroscience as a serotoninergic neurotoxin. However, consecutive studies regarding the mechanism of its action revealed the limitations of 5,6-DHT use (Baumgarten and Schlossberger 1973), which motivated scientists to utilize 5,7-dihydroxytryptamine (5,7-DHT), an isomer of 5,6-DHT.

# 2 Dihydroxytryptamines

### 2.1 Chemical Structure

Dihydroxytryptamines are indole derivatives that acquire their neurotoxicity due to additional hydroxyl groups (in comparison with 5-HT) on the indole ring (Schlossberger 1978).

#### 2.2 The Mode of Administration

5,6-DHT as well as 5,7-DHT can be administered intracerebroventricularly, intracerebrally, intracisternally, and intrathecally (Baumgarten et al. 1978b).

# 2.3 Cytochrome-c Oxidase and the Neurotoxicity of Dihydroxytryptamines

Cytochrome-c oxidase is a common feature in the mechanisms of toxicity of both 5,6-DHT and 5,7-DHT (Baumgarten et al. 1982b).

# 3 Dose-Dependent Action of Dihydroxytryptamines

#### 3.1 5,6-Dihydroxytryptamine

5.6-DHT was the first dihydroxytryptamine introduced to neurobiology. Its injection into the cerebrospinal fluid of rats at doses of 25–75 µg leads to degeneration of indolamine-containing axons and axon terminals in distinct regions of the brain and spinal cord (Baumgarten et al. 1971). At this range of doses, 5,6-DHT causes no considerable changes in DA and noradrenaline (NA) levels in any region of the rat brain (Baumgarten et al. 1972). The same doses injected intracerebroventricularly trigger long-lasting depletion of 5-HT in the spinal cord and smaller effects in the brain (Baumgarten et al. 1971; Daly et al. 1973). Doses higher than 75 ug do not enhance the neurotoxic effects of 5,6-DHT on serotonergic pathways and are extremely toxic to the animals, leading to muscle paralysis, convulsions, tremor, and death in the majority of animals. Furthermore, such doses cause damage to other types of neurons. Studies show nonselective destruction of myelinated fiber systems and glial cells and shrinkage of the striatum on the side of the injection. In doses above 75 µg administered intracerebroventricularly, 5,6-DHT induces a significant reduction of DA and NA levels in distinct areas of the brain (Baumgarten et al. 1972; Bjorklund et al. 1973). Bjorklund and collaborators (1973) demonstrated that this dose restriction can be overcome by infusion of small amounts of the neurotoxin directly into the brain parenchyma.

# 3.2 5,7-Dihydroxytryptamine

Limitations in the use of 5,6-DHT and need for neurotoxins with less general cytotoxicity inspired the discovery of more selective serotonergic neurotoxins. In 1972, Baumgarten and Lachenmayer described the effects of intracerebral injection of another 5-HT neurotoxin - 5,7-DHT. One of the basic distinctions between 5.6- and 5.7-DHT is the ability to use doses up to 300  $\mu$ g, which are ordinarily well tolerated. Further, the injection can be made intracerebroventricularly, so the neurotoxin does not have to be injected directly into brain tissue (Baumgarten and Lachenmayer 1972). However, doses up to 100 µg administered intracerebroventricularly in rats often lead to convulsions that can be prevented by previous treatment with barbiturates (Baumgarten and Lachenmayer 1972). Baumgarten and collaborators (1973) demonstrated that even 10  $\mu$ g of 5,7-DHT injected intracerebroventricularly triggers a significant reduction (approximately 25 %) in the levels of 5-HT and NA in the rat brain. A dose of 50 µg leads to a further decrease in the concentration both of neurotransmitters to approximately 50 % of control values. Successively increasing doses from 75 to 200 µg cause a further decline in the 5-HT level to about 25 % of the control value, without further reduction in the NA level. In early studies, even large doses of 5,7-DHT did not affect the brain levels of DA, a finding that was verified by subsequent research demonstrating that 5,7-DHT also damages DA axons (Wuttke et al. 1977; Baumgarten et al. 1978b). It is important that the neurotoxic action of 5,7-DHT is not equal in all central nervous system (CNS) regions. Doses as low as 10  $\mu$ g are sufficient to cause almost complete (85 %) reduction in the 5-HT level in the spinal cord (Baumgarten et al. 1973). Depletion in the 5-HT level in rats after injection of 200  $\mu$ g ranged from 55 % in the medulla and pons to approximately 85 % in the forebrain and septum (Baumgarten et al. 1973). The most considerable decrease in 5-HT level was noted in the regions of the CNS rich in serotonergic axons and terminals (i.e., the septum, striatum, and hypothalamus), whereas much lower depletion was observed in areas containing serotonergic cell bodies (i.e., mesencephalon, medulla/pons) (Baumgarten et al. 1973).

# 4 Dihydroxytryptamines: Differences

# 4.1 Neurotoxicity

A comparison of mono-, di-, and trihydroxylated tryptamines reveals that the principal factor affecting the neurotoxicity of these compounds is the velocity of their oxidation at biological pH. Moderate reaction velocity provides the best potential for toxicity. In the case of 5,6,7-trihydroxytryptamine, a very high oxidation rate causes its immediate transformation into a quinone derivative that does not serve as a substrate for monoaminergic transport mechanisms. A similar mechanism is responsible for the lower in vivo neurotoxicity of 5,6-DHT in comparison to 5,7-DHT, even though 5,6-DHT shows a higher affinity to the amine carrier of serotoninergic neurons (Baumgarten 1981, 1982b). 5,6-DHT undergoes rapid oxidation to o-quinone, which forms melanin-like polymers by aggregation and/or binds covalently and irreversibly to brain proteins (Baumgarten et al. 1975a). These features ensure that 5,6-DHT never penetrates into deeply located 5-HT uptake sites. Moreover, they reduce 5,6-DHT toxicity to 5-HT neurons and increase the risk of nonselective toxic side effects. Conversely, 5,7-DHT is not oxidized as easily, does not create insoluble polymers, and rapidly penetrates into the brain, thus giving it high intrinsic neurotoxicity despite a lower affinity for the 5-HT uptake mechanism (Baumgarten et al. 1975b).

The key feature that determines dihydroxytryptamines activity is the localization of the hydroxyl groups. 5,7-DHT is a meta-substituted dihydroxytryptamine, while 5,6-DHT is ortho-substituted. The meta-substitution enables 5,7-DHT to exist in solution as an equilibrium of distinct tautomeric keto-enol forms, which provide affinity to the 5-HT carrier despite the oxidation of some fraction of 5,7-DHT in vivo. Oxidation does not cause the total loss of 5,7-DHT affinity to the 5-HT carrier (unlike to 5,6-DHT) because one of the hydroxyl groups in the 5,7-DHT structure remains functional (Baumgarten et al. 1982b).

# 4.2 IC<sub>50</sub> for 5-HT Uptake Inhibition

IC<sub>50</sub> values for the inhibition of 5-HT uptake are 4  $\mu$ M for 5,7-DHT and 0.6  $\mu$ M for 5,6-DHT, making 5,7-DHT almost seven times less potent than 5,6-DHT (Bjorklund et al. 1975c). However, 5,7-DHT is a better long-term depletor of brain and spinal cord 5-HT than 5,6-DHT, possibly due to differences in metabolism and reactivity of 5,7-DHT (Baumgarten et al. 1975b).

#### 4.3 Selectivity for Catecholamine Uptake

Early studies suggested that 5,7-DHT was a superior serotonergic neurotoxin to 5,6-DHT due to lower nonspecific toxicity and higher specific neurotoxicity (Baumgarten et al. 1973), but further studies by Wuttke et al. (1977) and then by Baumgarten and collaborators (1978b) revealed that 5,7-DHT affects both central NA fibers and DA axons. The results by Bjorklund et al. (1975c) showed that 5,6-DHT is a more specific inhibitor of 5-HT uptake than catecholamine uptake, despite the small differences between the affinity of 5,6-DHT to the 5-HT, NA, and DA transporters (% inhibition of uptake is 59.0  $\pm$  2.3; 43.3  $\pm$  2.7; 44.7  $\pm$  0.7, respectively). 5.7-DHT has a different selectivity pattern. It inhibits the uptake of catecholamines, particularly NA, more effectively than 5-HT (% inhibition of uptake is  $25.7 \pm 1.2$ ;  $32.5 \pm 1.3$ ;  $15.4 \pm 1.7$  for 5-HT, NA, and DA, respectively (Bjorklund et al. 1975c)). These results are different from those presented by Baumgarten and collaborators (1978b). On the basis of K<sub>m</sub> values for NA uptake (10.0  $\times$  10<sup>-6</sup> M) and 5-HT uptake  $(6.0 \times 10^{-7} \text{ M})$ , it appears that 5,7-DHT has a 16-fold higher affinity for the 5-HT transport mechanism. However, there are also studies that show that 5,7-DHT exhibits dose-dependent differences in its selectivity. In doses up to 50  $\mu$ g, it depletes NA and 5-HT uptake to a similar degree, whereas in doses higher than 50  $\mu$ g, it affects CNS 5-HT levels substantially more (Baumgarten et al. 1971, 1973, 1975b).

The toxicity of dihydroxytryptamines to non-5-HT neurons can be counteracted by the use of nomifensine, which is a NA/DA uptake blocker (Baumgarten et al. 1979). The effects of 5,7-DHT on NA neurons can be abolished by desmethylimipramine, a blocker of the amine pump of NA neurons. In numerous studies, a combination of desmethylimipramine and 5,7-DHT has been used to achieve selective chemical ablation of 5-HT axons and terminals in the rat brain (Bjorklund et al. 1975a; Sachs and Jonsson 1975). Other methods to block the effect of 5,7-DHT on NA neurons include the use of monoamine oxidase inhibitors (MAOI) such as pargyline, iproniazid, and pheniprazine (Breese and Cooper 1975) and pretreatment with desipramine (Wuttke et al. 1977; Bjorklund et al. 1975b).

## 4.4 Half-Life and Stability

There are also differences between the half-life and stability of 5,6- and 5,7-DHT. 5,7-DHT is more stable because it lacks autocatalytic promotion of its oxidation

and does not create macromolecular polymers. In contrast, the auto-oxidation of 5,6-DHT is promoted by  $H_2O_2$ , and the product of its oxidation, o-quinone, reacts with proteins and peptides to generate high molecular weight polymers through interaction with non-oxidized 5,6-DHT (Baumgarten et al. 1982b).

### 4.5 The Rate of Oxygen Consumption

5,6- and 5,7-DHT differ in their rates of oxygen consumption. The initial  $O_2$  consumption is 2.7 nmol  $O_2$ /min for 5,6-DHT and 33.4 nmol  $O_2$ /min for 5,7-DHT. This difference reflects the fact that oxidation of 5,7-DHT is a second-order reaction. However, the rate of oxidation of 5,6-DHT increases after an initial period of slow oxygen consumption (Baumgarten et al. 1982b).

# 4.6 Diffusion

5,6-DHT has limited diffusion into the brain parenchyma of the adult rat due to its capacity to create macromolecular polymers (Baumgarten et al. 1977). 5,7-DHT infiltrates much deeper into the brain. Notably, 5,7-DHT has the ability to freely diffuse across the brain and to cross the blood-brain barrier during the first week of life. These features enable denervation of the entire 5-HT neuronal system following the intraperitoneal administration of 5,7-DHT to rat pups (Sachs and Jonsson 1975; Kostrzewa et al. 1994).

### 5 Factors That Affect the Action of 5-HT Neurotoxins

Factors influencing the action of dihydroxytryptamines include the site of injection, the speed and volume of injection, the amount and type of antioxidant used, the type of anesthesia, and the dose and type of catecholamine uptake blocker. These factors influence the specificity of the treatment, the extent of amine depletion, the time course of degeneration and regeneration, the development and recovery of a postsynaptic supersensitivity, and the symmetry of neurotoxic action (Baumgarten et al. 1982b).

### 5.1 The Site of Injection

The site of injection affects the degree and selectivity of the serotonin neurotoxin action, as demonstrated by comparing the effects of 5,7-DHT intracerebral injection into the lateral and fourth ventricle. Injection of 100  $\mu$ g of 5,7-DHT into the left lateral ventricle causes rapid loss of 5-HT from the forebrain with no considerable recovery up to 55 days following injection. Administration of 5,7-DHT into the fourth ventricle (termed intracisternal injection) evokes a less extensive and more delayed 5-HT depletion, with mild but significant recovery occurring

within 2 months of injection (Baumgarten et al. 1978a, b). The route of administration also influences the amount of nonspecific toxic damage to the CNS, which is higher when the neurotoxin is injected into the lateral ventricle than into fourth ventricle (Hedreen 1975).

# 5.2 The Speed and the Volume of Injection

The speed of 5,7-DHT injection affects the extent and symmetry of its action in periventricular structures when the neurotoxin is administered into one of the lateral ventricles. Comparison of two distinct velocities of injection revealed that slow infusion (1  $\mu$ l/min) causes a more extensive reduction of amine uptake and leads to a more symmetric loss of 5-HT uptake (75 % and 50 % in the left and right striatum, respectively), while fast injection (3 ul/min) leads to asymmetric decreases in 5-HT uptake (Gershanik et al. 1979). Furthermore, large interindividual discrepancies can result from different times elapsing between the neurotoxin administration and sacrifice (2–7 days). Time of sacrifice is an important factor because the damage of 5-HT axons is a time-dependent process (Baumgarten et al. 1973, 1977). Slow versus fast 5-HT neurotoxin administration also produces entirely different results when observed at different time points. One hour after infusion, rapid injection causes symmetric, whereas slow injection causes asymmetric reduction in the 5-HT content of the brain. However, when assessment of 5-HT levels is performed 24 h after neurotoxin administration, initial differences become equalized (Baumgarten et al. 1982b).

### 5.3 The Influence of Anesthesia

There are inconsistent results regarding the influence of anesthesia on the efficiency of 5,7-DHT neurotoxicity. Baumgarten and collaborators (1982a) suggest that administration of pentobarbitone with ketamine does not exert a major effect on 5,7-DHT action, whereas in an earlier study, the application of pentobarbitone attenuated the toxicity of 5,6-DHT (Baumgarten et al. 1972). Furthermore, the anesthetic can affect not only the neurotoxin activity but also the activity of the catecholamine uptake blocker that is used to gain the high selectivity of the neurotoxin action. Pentobarbitone was shown to interfere with nomifensine in animals given a slow injection of 5,7-DHT (Baumgarten et al. 1982a).

# 6 Mechanism of 5,6- and 5,7-Dihydroxytryptamine Neurotoxicity

The mechanism of action of dihydroxylated tryptamines is connected with their uptake into neuronal terminals by the active transport systems that are normally responsible for the reuptake of 5-HT (Baumgarten et al. 1978c). The neurotoxins accumulate

intraneuronally to achieve a critical concentration that causes axonal damage. The specificity and potency of each neurotoxin is determined by its affinity for a given monoamine uptake mechanism, its rate of clearance via cellular metabolism, and the ability of its breakdown products to bind to proteins (Bjorklund et al. 1975c). Two major hypotheses have been proposed to explain the molecular mechanism that underlies the toxicity of 5.6- and 5.7-DHT. The first suggests that highly reactive (electrophilic) quinonoid intermediates, created as a result of 5,6- and 5,7-DHT auto-oxidation, bind covalently to molecules indispensable for neuronal function (Rotman et al. 1976). The second hypothesis states that during the auto-oxidation of dihydroxylated tryptamines, reactive oxygen species (ROS) such as superoxide radical anion  $(O_2^{-})$ , hydrogen peroxide  $(H_2O_2)$ , and the hydroxyl radical (HO) are generated as toxic by-products (Baumgarten et al. 1978b, 1982a, b). Studies show that free radical scavengers can provide some protection to peripheral nerves against the degeneration caused by 5.7-DHT (Cohen and Heikkila 1978). Moreover, it has been speculated that quinine imine intermediates of 5,7-DHT can also trigger toxicity via an interaction with the electron transport chain (Klemm et al. 1980).

# 6.1 Mechanism of 5,7-DHT Action

The auto-oxidation of 5,7-DHT in a buffered aqueous solution at pH 7.4 causes the solution to turn pink (Schlossberger 1978; Cohen and Heikkila 1978; Tabatabaie et al. 1990). The auto-oxidation reaction is second order (Klemm et al. 1980) and at physiological pH leads to the formation of two major products – 5-hydroxytryptamine-4,7-dione and 6,6-bi-5-hydroxytryptamine-4,7-dione (Tabatabaie et al. 1990). Auto-oxidation of 5,7-DHT produces molecular oxygen that catalyzes the oxidation of brain mitochondria (Cohen and Heikkila 1978). This result was confirmed in 1980 by Klemm and collaborators. Moreover, they ruled out any pivotal participation of MAO in the catalytic action of mitochondria. Mitochondria play an important role in catalyzing the auto-oxidation of 5,7-DHT and may be sites of accumulation of damage caused by 5,7-DHT (Tabatabaie and Dryhurst 1998). In 1992, Tabatabaie and Dryhurst revealed that in the presence of superoxide dismutase (SOD), catalase, and diethylene triamine pentaacetic acid (DTPA), which eliminate the catalytic action of  $O_2^-$ , trace transition metal ions, and organic peroxy intermediates, the auto-oxidation of 5,7-DHT is much slower than normal. Among the transition metal ions, Fe<sup>3+</sup>, Fe<sup>2+</sup>, Cu<sup>2+</sup>, and Mn<sup>2+</sup> (Tabatabaie and Dryhurst 1992) are strong candidates to catalyze 5,7-DHT auto-oxidation. The rate of auto-oxidation decreases after catalase and SOD application but increases after the addition of  $O_2^-$ . The lack of a stimulatory effect after  $H_2O_2$  administration is striking, especially considering inhibitory catalase action. It has been suggested that the ability of catalase to decompose organic peroxides plays pivotal role in the auto-oxidation of 5,7-DHT (Kadlubar et al. 1973). O<sub>2</sub><sup>-</sup> also plays a role in autooxidation of 5,7-DHT. It is not only a by-product, but it also plays significant role in potentiating the rate of the reaction. Studies suggest that the catalytic action of  $O_2^{-1}$ is not directly connected with oxidation of 5,7-DHT because  $O_2^-$  is a weak

oxidizing agent (Sawyer et al. 1978). However, it may react with organic hydroperoxides and contribute to a free radical chain reaction (Tabatabaie and Dryhurst 1998). Moreover, it has been proposed that  $O_2^-$  stimulates the deprotonation of 5,7-DHT, which is the initial step in auto-oxidation (Tabatabaie and Dryhurst 1992).

Due to its structure similarity to 5-HT, 5.7-DHT is taken up and concentrated in serotonergic nerve terminals (Baumgarten et al. 1978c). There, the result of 5,7-DHT oxidation is the generation of 5-hydroxytryptamine-4,7-dione and 6,6-bi-5-hydroxytryptamine-4,7-dione as major products. Both of these compounds are more toxic than 5,7-DHT. Their LD<sub>50</sub> values, measured by ventricular administration in mice, are 30 and 25  $\mu$ g, respectively, whereas the LD<sub>50</sub> for 5,7-DHT is 52  $\mu$ g (Tabatabaie et al. 1993). However, they do not have the ability to cause profound and long-lasting depletion of 5-HT as occurs after 5,7-DHT administration (Tabatabaie and Dryhurst 1998). The result of 5,7-DHT oxidation is the formation of  $O_2^-$  and  $H_2O_2$ , which, in the presence of trace levels of transition metal ions, cause the generation of cytotoxic HO' (Halliwell 1992). The other effect of this auto-oxidation is increased consumption of molecular oxygen, leading to hypoxia, changes in metabolism, and cellular damage (Kappus 1986). The auto-oxidation product 5-hydroxytryptamine-4,7-dione also reacts with glutathione (GSH) to cause depletion of GSH storage, exacerbating damage caused by ROS (Tabatabaie and Dryhurst 1998).

# 6.2 Mechanism of 5,6-DHT Action

The ultimate product of 5,6-DHT auto-oxidation at physiological pH is a black, insoluble, melanin-like polymer (Klemm et al. 1980). The by-product of this reaction is a high amount of  $H_2O_2$  (Klemm et al. 1980; Singh and Dryhurst 1990). Studies show that  $H_2O_2$ ,  $O_2^-$ , and trace levels of transition metal ions play an important role in 5,6-DHT auto-oxidation because the application of catalase, SOD, and iron-complexing agents inhibit this reaction. However, the removal of agents such as transition metal ions and molecular oxygen from the reaction site causes 5,6-DHT not to be oxidized by  $H_2O_2$  (Singh and Dryhurst 1990). In turn, the application of even very low amounts of  $Fe^{3+}$ ,  $Cu^{2+}$ , and  $O_2^{-}$  augments the oxidation of 5,6-DHT. These results indicate that  $H_2O_2$  does not participate directly in 5,6-DHT auto-oxidation, but it undergoes a transformation (via the transition metal ion-catalyzed Fenton reaction) leading to HO generation, which directly affects 5,6-DHT. In vitro as well as in vivo studies show that products of 5,6-DHT auto-oxidation undergo covalent binding with protein nucleophiles (Baumgarten et al. 1978b). However, auto-oxidation of 5,6-DHT ultimately leads to melanin-like polymer formation. An assessment of the role of covalent binding products of 5,6-DHT auto-oxidation revealed that they are not essential for the cytotoxic effects of 5,6-DHT (Sinhababu et al. 1985). The other mechanism that triggers the neurodegenerative action of 5,6-DHT is its ability to deplete intraneuronal antioxidants (e.g., GSH) and generate ROS that induce oxidative damage of lipids, proteins, and

DNA (Tabatabaie and Dryhurst 1998). The result of intraneuronal auto-oxidation of 5,6-DHT is the alkylation of proteins and, in favorable conditions, the cross-linking of proteins with GSH. Both reactions can lead to lethal consequences (Tabatabaie and Dryhurst 1998).

# 7 Behavioral Effects

At the beginning of this subsection, we would like to emphasize that there are agerelated differences in the response of 5-HT neurons to dihydroxytryptamines. Young neurons display increased sensitivity to dihydroxytryptamines due to underdeveloped detoxifying mechanisms such as degrading enzymes, 5,7-DHT storage capacity, and the ability to scavenge radicals and neutralize oxidation intermediates through nucleophilic compounds such as GSH (Breese and Mueller 1978). Moreover, the accumulation of 5,7-DHT in the relatively small 5-HT perikarya of neonatal rats is more efficient than in adult rats where the surface/volume relationship is less favorable. Brain weight has to be taken into account because the brain weight of adult rats is about tenfold higher than newborn rats (Baumgarten et al. 1982b). On the other hand, infant brains have a greater potential for functional recovery than adult brains because they have greater potential for neuronal regeneration (Jonsson et al. 1978).

# 7.1 Adult Animal Studies

#### 7.1.1 Changes in Locomotor Activity

Intraventricular administration of one of the dihydroxytryptamines (5,6- or 5,7-DHT) to rats causes dramatic behavioral changes (Massotti et al. 1974). Behavioral depression and sedation during the first 2 h after the infusion is typical (Baumgarten and Schlossberger 1973; Massotti et al. 1974). Following this initial period, increased irritability, explosive running, disorganized locomotion, and hyperreactivity to any type of sensory stimulation are observed. These responses are accompanied by bizarre social behavior (fighting in a stereotyped position and manner) and increased copulatory activity between the 1st and 3rd day after 5,6-DHT administration (Baumgarten and Schlossberger 1973). The greatest locomotor hyperactivity occurs during the first week after neurotoxin (5,7-DHT) administration (Pranzatelli and Snodgrass 1986b), and increased locomotor activity (LMA) persists 1-3 weeks following the intracisternal (Stewart et al. 1979) and intracerebroventricular (Lyness and Moore 1981) injection of 5,7-DHT. The stimulation of LMA has been observed after stereotactic injection of 5,7-DHT into the nucleus accumbens septi, substantia nigra, and striatum (Lyness and Moore 1981; Carter and Pycock 1981). The destruction of the medial raphe nuclei (MRN) also leads to increased LMA that manifests mainly as nocturnal hyperactivity and is correlated with hippocampal 5-HT depletion and a limbic mechanism (Geyer et al. 1976). The forebrain may be involved in the changes to LMA following the serotonergic manipulations (Geyer et al. 1976; Carter and Pycock 1981).

After this period of increased LMA, behavior returns to normal, and it is difficult to distinguish DHT-treated animals from controls. This normalization is termed "recovery" and is related to functional reorganization and supersensitivity (Glick 1974; Finger and Almli 1985). Despite the recovery of LMA, the supersensitivity to 5-hydroxy-L-tryptophan (L-5-HTP) may remain for at least 12 months after 5,7-DHT administration (Pranzatelli and Snodgrass 1986a). In 1978, Mackenzie and collaborators revealed that serotonin depletion by 5,7-DHT in rats causes tilt-cage hyperactivity but decreases activity in the open-field test. Kusljic et al. (2003) showed differential involvement of serotonergic projections deriving from medial raphe nuclei and dorsal raphe nuclei (DRN) in the regulation of locomotor hyperactivity. 5,7-DHT injected into the MRN causes a marked increase of phencyclidine-induced locomotor hyperactivity, whereas administration of DHT into the DRN does not.

#### 7.1.2 Prepulse Inhibition

Prepulse inhibition (PPI) is a normal reduction of a startle response taking place when a weak stimulus occurs within a very short time period before a startleevoking stimulus (Graham 1975). PPI measures sensorimotor gating and reflects the ability to filter out irrelevant stimuli. Fletcher et al. (2001) found that severe depletion of 5-HT (by 5,7-DHT injection into the DRN and MRN) disrupts PPI of the acoustic startle reflex without influencing basal startle activity. Kusljic and coworkers (2003) demonstrated that, similar to LMA, the MRN and DRN regulate PPI in different ways. 5,7-DHT injected into the MRN or the DRN leads to disruption of PPI, but the PPI effect is more significant in MRN-lesioned rats, while the startle amplitude is markedly increased in DRN-lesioned animals.

#### 7.1.3 Seizure Susceptibility

The contribution of serotonin to seizure susceptibility was first suggested in 1954 by Chen and collaborators (Chen et al. 1954). They revealed that the seizure-facilitating effects of reserpine derive from its depletion of 5-HT as well as catecholamines. In adult rats pretreated with protriptyline and 5,7-DHT, the severity of pentylenetetrazol (PTZ)-induced seizures is augmented. These animals also display an earlier seizure onset and a greater severity of seizures than control rats that received vehicle pretreatment. Thus, 5,7-DHT-treated rats seem to be more seizure prone than controls (Browning et al. 1978). Similar results have been obtained by Trindade-Filho and coworkers (2008), who assessed the effects of 5,7-DHT administration into the MRN in a rat pilocarpine model of epilepsy. They showed that the frequency of seizures in treated animals was significantly increased compared to control rats.

#### 7.1.4 Serotonin Syndrome

The classical 5-HT syndrome manifests as lateral head "weaving" (side-to-side movement), reciprocal forepaw "treading," "resting tremor," hindlimb abduction, rigidly arched (Straub) tail, hyperreactivity to auditory stimuli, limb and axial rigidity or hypertonicity, and autonomic responses (salivation, erection, and

ejaculation) (Jacobs 1976). Dihydroxytryptamines evoke supersensitivity to 5-HT precursors and agonists. Studies in 5,7-DHT-treated adult rats revealed that application of L-5-HTP (precursor of 5-HT) leads to 5-HT syndrome (Browning et al. 1978). Low doses of 5-hydroxytryptophan (5-HTP) cause a self-limiting syndrome characterized by the appearance of myoclonus, other dyskinesias, stereotypies, and abnormalities of posture and tone (Stewart et al. 1976, 1979; Pranzatelli and Snodgrass 1986b). It has been suggested that the brainstem and spinal cord are the brain areas involved in mediating this serotonergic behavioral syndrome (Jacobs and Klemfuss 1975).

### 7.1.5 Feeding and Drinking Behavior

Intraventricular injection of dihydroxytryptamines leads to immediate and transient anorexia, hypodipsia, and weight loss. However, treated animals quickly return to normal body weight and show normal patterns of food and water intake; permanent alteration of drinking preference is rare (Myers 1975; Melchior and Myers 1976). Saller and Stricker (1976) suggested that the inability to observe consistent changes in digestive behavior derives from the effect of dihydroxytryptamines on neurons containing NA. They showed that animals treated with desmethylimipramine and subsequently intraventricularly with 5,7-DHT become hyperphagic and obese, but animals injected with only 5,7-DHT show no change or minor reductions in eating, drinking, and body weight (Saller and Stricker 1976). Opposite results were reported by Myers (1978) in monkeys and rats treated with 5,6-DHT infused directly into hypothalamic structures. A significant reduction of food consumption was observed in monkeys following neurotoxin infusion in the anterior hypothalamic, preoptic region. Water consumption was decreased but not significantly different from the baseline intake. In most cases, the decline of food and water intake occurred not only on the day of 5.6-DHT injection but also on the following day. In rats, hypophagia lasts 4–5 days after the infusion. Water consumption in rats was changed in a similar manner as was food intake. Further, rats lost between 30 g and 50 g of body weight at the higher doses of 5,6-DHT (Myers 1978). The comparison of food intake and body weight studies involving administration of 5,6-DHT, 5,7-DHT, and 6-OHDA to different strains of rats reveals that 5,7-DHT injected intraventricularly produces the greatest loss of body weight (Melchior and Myers 1976).

#### 7.1.6 Body Temperature

In 1963, it was first suggested that 5-HT within the anterior hypothalamus participates in the control of heat production (Feldberg and Myers 1963). Currently, we know that neurons containing 5-HT take part in the regulation of body temperature, but we still do not know the exact role of this transmitter in the mechanism due to inconsistent results between different studies. The effects of dihydroxytryptamines depend on the species and the interval between drug injection and examination. In rats, intraventricular injection of 5,6-DHT causes immediate, transient hypothermia (Breese et al. 1974a). On the other hand, studies in rats and monkeys suggest that infusion of 5,6-DHT into the anterior hypothalamus causes not hypo- but hyperthermia (Waller and Myers 1974; Myers 1975). Myers (1978) emphasizes that the thermic response to 5,6-DHT is dose dependent and if too large a dose of neurotoxin is injected, transient hypothermia always precedes the hyperthermia.

#### 7.1.7 Sexual Behavior

Neurons containing 5-HT mediate the inhibition of copulatory behavior (Meyerson 1964, 1966). Studies using serotonergic neurotoxins confirm that serotonin nerves inhibit sexual behavior and depletion of 5-HT increases sexual activity in rats (Da Prada et al. 1972). Intracerebroventricular injection of 5,6-DHT induces hyperexcitability and fighting in rats. The main characteristic effect of such treatment is mounting activity (Da Prada et al. 1972; Baumgarten and Schlossberger 1973). If 5-HT levels are normalized by 5-HTP administration, mounting behavior is completely abolished. Similar effects were observed in 5,7-DHT-lesioned rats treated with 5-HTP. Grouped female rats injected with 5,7-DHT also exhibit sexual activity analogous to the mounting displayed by male rats. A similar effect can be triggered by 5,6-DHT treatment. In addition, protection of 5-HT neurons with the 5-HT inhibitor Lu 10-171 before 5,7-DHT administration prevents fighting and mounting behaviors (Da Prada et al. 1978).

### 7.1.8 Aggressive Behavior

Serotonergic neurons may play a pivotal role in predatory and irritable aggression, defined as aggressive behavior against an animal of another species and against animals of any species triggered by an aversive stimulus, respectively (Eichelman and Thoa 1973). Depletion of 5-HT increases predatory aggression toward mice (also termed muricidal behavior). Intracisternal injection of 5,6-DHT evokes a marked increase in muricidal behavior, even 1 week after infusion. Pretreatment with pargyline significantly augments this predatory aggression. Similar results were found in rats injected with 5,6-DHT in the midbrain raphe nuclei. However, pargyline pretreatment did not significantly increase muricidal behavior, observed 19 days after neurotoxin administration. Treatment with 5,7-DHT, pargyline + 5,7-DHT, or desipramine + 5,7-DHT causes killing in 71 %, 89 % and 40 % of rats, respectively (Breese and Cooper 1975).

#### 7.1.9 Response to Novel and Noxious Stimuli

The presentation of a novel stimulus triggers exploratory activity and behavioral vigilance. Subsequent exposures to the same stimulus cause a graded attenuation of these behaviors until they completely disappear. Neural processes termed habituation and sensitization are responsible for this behavior (Davis and Sheard 1974). 5-HT is involved in both the sensitization process (Carlton and Advokat 1973) and the response to a noxious stimulus (Messing and Lytle 1977). However, investigations with 5,6-DHT or 5,7-DHT injected intraventricularly (Blasig et al. 1973) or directly into the midbrain (Hole and Lorens 1975) did not reveal any changes in the behavioral response to noxious stimuli. Lieben and collaborators (2006) found that 5,7-DHT lesion of the dorsal raphe nuclei impairs the ability to identify a novel object.

#### 7.1.10 Learning and Memory Functions

It has been suggested 5-HT participates in both short-term and long-term memory (Bailey et al. 1992; Barbas et al. 2003). There is some literature showing that 5,7-DHT injection does not change learning and memory capabilities even when severe 5-HT depletion is induced electrolytically (Asin and Fibiger 1984). However, there are also data providing quite distinct results that show that 5,7-DHT administration affects short-term memory. Hritcu and collaborators (2007) found that disruption of short-term memory does not accompany alterations in long-term memory. Ciobica et al. (2010) found a significant defect in short-term memory, demonstrated by working memory errors in the radial arm maze. Ricaurte and coworkers (1993) noted an impairment of choice accuracy following 5,7-DHT treatment, which also suggests alterations in short-term memory. Moreover, Cassaday et al. (2003) found that rats treated with 5,7-DHT had difficulty in learning new tasks but no impairment in performing previously learned working memory tasks. However, data demonstrating a lack of short-term memory deficits after neurotoxin injection also exist (Wirth et al. 2000).

## 7.1.11 Effect of Dihydroxytryptamines on the Antinociceptive Actions of Drugs

5-HT is involved in determining the pain threshold and in regulating the perceptual and behavioral responses to painful stimuli (Lorens 1978). Decreasing 5-HT levels (by different methods) reduces the analgesic potency of morphine (Tenen 1968), whereas increasing 5-HT concentrations augments morphine analgesia (Sewell and Spencer 1974). However, not all results confirm this hypothesis. Some studies indicate that mesencephalic raphe lesions and intra-midbrain infusion of 5,7-DHT (10 µg) do not change pain sensitivity (Harvey et al. 1974; Hole and Lorens 1975). Blasig and collaborators (1973) noticed that intraventricular injection of 5,6-DHT did not significantly change the antinociceptive effect of morphine. Moreover, they showed that neurotoxin did not affect the stimulation threshold before morphine administration. These results suggest that a decreased 5-HT level is not sufficient to trigger the development of hyperalgesia (Lorens 1978). On the other hand, the majority of studies show different results. Genovese and collaborators (1973) found that, following intraventricular injection of 5,6-DHT in rats, the pain threshold was modestly but not significantly reduced. Following morphine infusion, the pain threshold was clearly increased in controls and sham-injected animals, but in rats that received 5,6-DHT, it was only slightly augmented. This result demonstrates that 5,6-DHT treatment reduces the antinociceptive action of morphine. Similar effects have been obtained in intrathecally 5,6- and 5,7-DHT-injected rats. Both neurotoxins evoke a decrease in the analgesic action of morphine, and no marked distinctions were noted between animals treated with 5,6- or 5,7-DHT (Rodriguez and Rodriguez 1989).

### 7.1.12 Sleeping Behavior

Manipulations that reduce forebrain 5-HT also decrease the duration of slow-wave sleep. This suggests a pivotal role for 5-HT in the initiation as well as maintenance

of this stage of sleep (Jouvet 1969, 1972). Studies in cats have shown that intraventricular injection of 5.6-DHT causes long-term reductions in slow-wave and REM (rapid eve movement) sleep time (Jouvet and Pujol 1974). Analogous results were not observed following intraventricular infusion of 5.7-DHT in rats. 5.7-DHT does not prominently affect the circadian pattern of sleep. The interesting difference between 5,7-DHT-treated and control rats was the amount of paradoxical sleep, which was significantly higher for the 5,7-DHT group on days 6 and 8. Additionally, 5,7-DHT rats displayed a large number of long duration paradoxical sleep epochs (Ross et al. 1976). Moreover, Jouvet and Pujol (1974) showed that direct administration of 5,6-DHT into the dorsal raphe or superior centralis nuclei of cats leads to a transient decrease in slow-wave sleep and markedly increases the duration of REM sleep within the first 24 h after treatment. These parameters subsequently return to normal. In 1978, Pujol and collaborators revealed that selective and diffuse damage of 5-HT terminals in cats causes a marked increase in waking time. Such evoked insomnia might be derived from decreases of stage 2 and paradoxical sleep as well as a simultaneous increase of the waking mechanism.

# 7.1.13 Neuroendocrine Consequences of Dihydroxytryptamine Administration

Many studies of the role of 5-HT in neuroendocrine regulatory mechanisms have used dihydroxytryptamines. In 1974, Ladosky and Noronha (1974) revealed that 5,6-DHT prominently increases the release of gonadotropin. However, other studies have provided different results. Significant changes in plasma and pituitary follicle-stimulating hormone (FSH) levels were not observed in rats treated at 21–23 days of age with 5,6-DHT (Saller and Stricker 1976). The influence of 5,7-DHT on plasma prolactin concentration has also been studied. Gil-ad et al. (1976) demonstrated that 5,7-DHT causes a reduction in prolactin levels, whereas Wuttke et al. (1978) and Clemens (1978) found no impairments in prolactin release in animals injected with 5,7-DHT. There are also inconsistent results in studies of the influence of 5-HT neurotoxins on growth hormone (GH). Collu (1978) showed that 5,6-DHT does not affect GH levels in male prepubertal rats, whereas Baumgarten and Bjorklund (1976) found that administration of 5,7-DHT increases plasma GH concentration.

#### 7.1.14 Effect of 5,7-DHT on the Dopaminergic System

When 5,7-DHT was administered in adulthood to rats that were neonatally lesioned with 6-hydroxydopamine, attenuated dopamine  $D_1$  receptor supersensitivity and intense spontaneous hyperlocomotor activity were observed (Kostrzewa et al. 1994). Amphetamine and m-chlorophenylpiperazine attenuation of the profound hyperlocomotion in rats lesioned with both 6-hydroxydopamine and 5,7-DHT reinforces the suggestion that such rats are a useful animal model of attention deficit hyperactivity disorder (Kostrzewa et al. 1994; Brus et al. 2004; Nowak et al. 2007).

# 7.2 Neonatal Animal Studies

# 7.2.1 Locomotor Activity

Studies suggest that serotonergic pathways play an inhibitory role in LMA (Breese et al. 1974b; Hollister et al. 1976). Neonatal rats injected with 5,7-DHT also display hyperactivity that persists for 1 month after the injection of neurotoxin. Breese and Mueller (1978) showed that the depletion of brain amines was greater in juveniles than in adults despite the lower doses of 5,7-DHT used. Moreover, they reported a significant fivefold elevation in LMA at 14 days in animals treated with 50  $\mu$ g of 5,7-DHT at 3 days of age. The LMA of animals treated with pargyline prior to 5,7-DHT was comparable to rats treated 5,7-DHT alone. This result suggests that rats pup hyperactivity does not require a reduction of NA fibers that increased LMA in adult rats (Smith et al. 1973). However, when the rats are tested at 28 days of age, the neurotoxin-treated groups demonstrate considerably less LMA than the control group (Breese and Mueller 1978).

# 7.2.2 Seizure Susceptibility

Pranzatelli and collaborators (Pranzatelli et al. 1989) demonstrated that intracisternal application of 5,7-DHT causes transient postinjection convulsions in rat pups. The first infusion leads to continuous paddling and writhing for at least 30 min without forward locomotion or vocalization. Moreover, 11 of 13 rats that received 5,7-DHT were unable to stand upright and fell over on their backs or sides. Subsequent 5,7-DHT injections caused more severe convulsions, whereas intraperitoneal injection of 5,7-DHT did not induce seizures during the 2-hour period of observation (Pranzatelli et al. 1989).

# 7.2.3 Serotonin Syndrome

Intracisternal and intraperitoneal injection of 5,7-DHT into rat pups, followed by the administration of 5-HTP 2 and 14 weeks later, leads to behavioral supersensitivity in adult rats. Both routes of 5,7-DHT administration provide supersensitivity to 5-HTP, but shaking behavior is greater following intraperitoneal infusion, whereas forepaw myoclonus and head weaving are greater following intracisternal injection (Pranzatelli et al. 1989).

# 7.2.4 Body Weights

A comparison of body weights between 5,7-DHT-treated rats and control animals at 14 days of age revealed that the former had markedly lower body weights. Pretreatment with pargyline prevents this weight loss. One possible explanation of this difference is that neurotoxin-treated rat pups bite the mother during suckling, which causes them to receive less nutrition (Breese and Mueller 1978).

# 7.2.5 Learning and Memory Function

Volpe et al. (1992), in their research with intracisternally administered 5,7-DHT to three-day-old rat pups pretreated with pargyline, revealed that 5,7-DHT destroys 5-HT neurons but does not influence spatial memory, tested by the Morris water

maze and the 12-arm radial maze. We also demonstrated that despite a severe and permanent decrease in hippocampal, prefrontal, and striatal 5-HT levels, neonatal treatment with 5,7-DHT caused no spatial learning and memory impairment in the Morris water maze (Piechal et al. 2012).

#### 7.2.6 Emotional and Exploratory Behaviors

In our study, neonatal 5,7-DHT depletion reduced LMA in the open-field test and attenuated social interaction in non-aversive conditions and exploration of a novel objects in adult rats. Ultravocalization, but not freezing, was increased in the contextual fear-conditioning paradigm in 5-HT-depleted rats. There was no difference in the pain threshold between groups (Rok-Bujko et al. 2012).

#### 7.2.7 Prepulse Inhibition

Our results clearly demonstrated that neonatal 5,7-DHT depletion does not interrupt PPI of the acoustic startle reflex. In contrast to the effect on PPI, neonatal treatment with 5,7-DHT increased basal startle activity (Kołomańska et al. 2011).

#### 7.2.8 Developmental Plasticity

Studies using several doses and schedules of 5,6-DHT administration in rat pups revealed that this neurotoxin is not useful for research in developmental studies because most pups died soon after treatment. Conversely, 5,7-DHT is well tolerated by young animals, even when injected repeatedly in doses up to 200 mg/kg. Systemic injection of 5,7-DHT in neonatal rats caused changes in 5-HT neurons that did not occur in adult animals. These changes include significant denervations of most distally located 5-HT nerve terminals, but cell bodies often remain intact. Moreover, in regions adjacent to cell bodies, compensatory outgrowth occurs that causes an increased number of 5-HT nerve terminals. This mechanism leads to redistribution of the terminals and bifurcations of 5-HT neurons and is termed "pruning" and "sprouting," respectively (Jonsson et al. 1978).

# 7.2.9 Effect of 5,7-DHT on Dopaminergic, Noradrenergic, and Histaminergic Systems

Neonatal 6-hydroxydopamine treatment is associated with supersensitization of dopamine  $D_1$  receptors, and this effect was attenuated by neonatal 5,7-DHT depletion (Brus et al. 1994). In contrast, the neonatal destruction of 5-HT fibers by 5,7-DHT produces supersensitization of the dopamine  $D_2$  receptor complex (Brus et al. 1995). Furthermore, the ontogenic destruction of serotonergic and noradrenergic neurons in rat brain is associated with prominent  $D_2$  and  $D_3$  receptor supersensitivity, as reflected by enhanced dopamine  $D_2$  and  $D_3$  agonist-induced stereotyped activities (Nowak et al. 2009). These findings indicate that both serotonergic and noradrenergic fibers have important influence on the process of dopamine  $D_1$  and  $D_2$  receptor sensitivity.

Finally, there is some evidence that ontogenetic 5-HT depletion alters histaminergic activity in adult rats. Neonatal 5,7-DHT treatment of rats was associated with enhanced oral and locomotor activity induced by the histamine  $H_3$  receptor antagonist thioperamide (Jośko et al. 2011).

### 8 Conclusion

5.6-dihydroxytryptamine (5.6-DHT) and 5.7-dihydroxytryptamine (5.7-DHT) are serotonin neurotoxins able to selectively destroy central serotonin neurons when catecholaminergic neurons are protected. Both compounds are indole derivatives, and their neurotoxicity derives from the additional hydroxyl group compared to serotonin (5-HT). The administration of 5,6- or 5,7-DHT to animals is a useful method to study the role of serotonin neurons and pathways in the central nervous system (CNS). Despite high structural similarity, 5,6-DHT differs from 5,7-DHT. The basic differences are connected to the ability of 5,6-DHT to rapidly oxidize and form consecutive melanin-like polymers. These features lower the neurotoxicity of 5.6-DHT to 5-HT neurons and make it more generally cytotoxic. Therefore, 5,7-DHT is used more frequently. The action of serotonin neurotoxins depends on the proper selection of injection sites, the speed and volume of administration. the type and amount of antioxidant used, and the type and dose of catecholamine uptake blocker used to prevent nonspecific toxicity. The mechanism of dihydroxvtryptamine action assumes that, because of their structurally similarity to 5-HT, they accumulate in 5-HT neurons where the products of their oxidation bind covalently to molecules indispensable for neuronal function. Moreover, reactive oxygen species (ROS) generated as by-products of oxidation also contribute to the destruction of 5-HT neurons. The use of 5,6- and 5,7-DHT in both adult and neonatal animals enables the recognition the role of 5-HT in locomotor activity (LMA), prepulse inhibition, seizure susceptibility, sleeping behavior, feeding and drinking behavior, sexual and aggressive behavior, the regulation of body temperature, the response to novel and noxious stimuli, learning and memory, the antinociceptive action of drugs, neuroendocrine regulatory mechanisms, and developmental plasticity.

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# 2'-NH<sub>2</sub>-MPTP: A Serotonin and Norepinephrine Neurotoxin

# Jason B. Ochroch, Amanda J. Bressler, Hongyan Yang, Dennis L. Murphy, Stefanie C. Altieri, and Anne M. Andrews

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### Abstract

Altered serotonin and norepinephrine neurotransmission has been implicated in a variety of psychiatric and neurodegenerative diseases. Selective ablation of these systems represents one avenue for studying their roles in normal behavior, disease processes, and therapeutic interventions. 2'-NH<sub>2</sub>-MPTP, an amine-substituted analog of the dopaminergic neurotoxin MPTP, can be used to degenerate serotonin and/or norepinephrine forebrain innervation to investigate

D.L. Murphy

e-mail: DennisMurphy@mail.nih.gov

J.B. Ochroch • A.J. Bressler • H. Yang • S.C. Altieri • A.M. Andrews

David Geffen School of Medicine, Department of Chemistry and Biochemistry, Semel Institute for Neuroscience and Human Behavior, and Hatos Center for Neuropharmacology, University of California, Los Angeles, CA, USA

e-mail: Jason.Ochroch@gmail.com; amandabressler75@gmail.com; yanghy@mednet.ucla.edu; stefaniealtieri@gmail.com; anne.andrews@ucla.edu

Laboratory of Clinical Science, National Institute of Mental Health, National Institutes of Health, Bethesda, MD, USA

pathophysiological and behavioral consequences in several strains of mice, rats, and nonhuman primates. 2'-NH<sub>2</sub>-MPTP is converted to a toxic metabolite by monoamine oxidase type-A and type-B. This metabolite, a presumed pyridinium, is a substrate for serotonin and norepinephrine transporters. Acute effects immediately following 2'-NH<sub>2</sub>-MPTP administration are characteristic of serotonin syndrome. Genetic, biochemical, and immunocytochemical evidence indicates that neurodegeneration caused by 2'-NH<sub>2</sub>-MPTP is selective, long lasting, and occurs by an oxyradical mechanism.

#### Keywords

Mice • Monkeys • MPTP analog • Neurodegeneration • Norepinephrine • Rats • Serotonin

List	of	Abbreviations	

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$2'-NH_2-MPP^+$	1-Methyl-4-(2'-aminophenyl)-phenylpyridinium			
$2'-NH_2-MPTP$	1-Methyl-4-(2'-aminophenyl)-1,2,3,6-tetrahydropyridine			
5-HIAA	5-Hydroxyindoleacetic acid			
5-HT	5-Hydroxytryptamine serotonin			
5,6-DHT	5,6-Dihydroxytryptamine			
5,7-DHT	5,7-Dihydroxytryptamine			
BDNF	Brain-derived neurotrophic factor			
CSF	Cerebrospinal fluid			
DA	Dopamine			
DAT	Dopamine transporter			
DOPAC	3,4Dihydroxyphenylacetic acid			
DSP-4 N	(2-Chloroethyl)-N-ethyl-2-bromobenzylamine hydrochloride			
GFAP	Glial fibrillary acidic protein			
HVA	Homovanillic acid			
Ip	Intraperitoneal			
Iv	Intravenous			
LD <sub>50</sub>	Median lethal dose			
MAO-A	Monoamine oxidase type-A			
MAO-B	Monoamine oxidase type-B			
MHPG	3-Methoxy-4-hydroxyphenylglycol			
$MPP^+$	1-Methyl-4-phenylpyridinium			
MPTP	1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine			
NE	Norepinephrine			
NET	Norepinephrine transporter			
PET	Positron emission tomography			
Ro 4–1284	2-Hydroxy-2-ethyl-3-isobutyl-9,10-dimethoxy-1,2,3,4,5,6,7-			
	hexahydrobenzo[a]chinolizine			
SERT	Serotonin transporter			
VMAT2	Vesicular monoamine transporter type-2			

#### 1 Introduction

Changes in brain monoamine neurotransmitters, i.e., serotonin, norepinephrine, and dopamine, are involved in many neuropsychiatric illnesses and their treatment, including major depressive disorder, anxiety disorders, and schizophrenia (Nestler et al. 2008; López-Muñoz and Álamo 2012). Moreover, degeneration of monoamine neurotransmitter systems is linked with normal aging and late-onset neurological diseases such as Parkinson's disease and Alzheimer's disease. However, studying brain disorders is inherently complicated by difficulties associated with the long time frames over which pathologies develop, the relative inaccessibility of investigating neurotransmission in humans, modeling and relating complex pathophysiology to specific endophenotypes, and a poor understanding of underlying etiologies. As such, chemical neurotoxins that cause controlled and selective damage to monoamine neurotransmitter systems in animals provide avenues to investigate aspects of specific neurotransmitter loss and neurodegeneration as they pertain to brain disorders.

## 2 Discovery of 2'-NH<sub>2</sub>-MPTP

The neurotoxin 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) produces degeneration of dopamine (DA) nigrostriatal neurons, which is the hallmark of idiopathic Parkinson's disease. MPTP came to the forefront in the early 1980s when a group of northern California drug users developed irreversible parkinsonian-like symptoms (Langston and Palfreman 1995). MPTP, a by-product of the illicit synthesis of a derivative of the opioid meperidine, was discovered as the causative agent.

Many analogs of MPTP have been synthesized for structure-activity studies (Bradbury et al. 1985; Youngster et al. 1987, 1989; Heikkila et al. 1988; Booth et al. 1989; Johnson et al. 1989; Maret et al. 1990; Saporito et al. 1992). MPTP-related compounds with a substituent at the 2' position, i.e., 2'-R-MPTP,  $R = CH_3$ ,  $CH_2CH_3$ , Cl, F, and CF<sub>3</sub>, have been shown to be potent dopamine neurotoxins. Amine-functionalized MPTP analogs were of interest because of the ability to covalently cross-link primary amines to brain tissue via aldehyde fixation. When coupled with immunohistochemical detection, amine-substituted MPTP compounds could be useful in mapping distribution at the ultrastructural level (Johannessen et al. 1987). However, adding an amine substituent to various positions on the phenyl ring of MPTP led to analogs that were relatively weak dopamine neurotoxins in mice.

Interestingly, the 2'-amino-substituted MPTP derivative, 1-methyl-4-(2'-aminophenyl)-1,2,3,6-tetrahydropyridine (2'-NH<sub>2</sub>-MPTP; Fig. 1a), produced reductions in striatal serotonin (5-HT) and its metabolite, 5-hydroxyindoleacetic acid (5-HIAA), while having no effect on dopamine in striatum (Johannessen et al. 1987). Because the goals of Johannessen et al. were to use amino-substituted analogs of MPTP to study toxicity related to Parkinsonism, no further work with

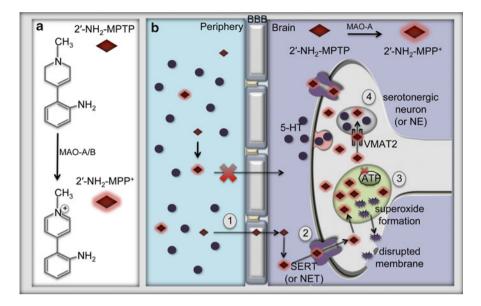


Fig. 1 Mechanisms of serotonin and norepinephrine neurotoxicity after exposure to 2'-NH<sub>2</sub>-*MPTP*. (a) Shown is the chemical structure of 2'-NH<sub>2</sub>-MPTP, a structural analog of the dopamine neurotoxin MPTP. Toxicity due to 2'-NH<sub>2</sub>-MPTP is inhibited by pretreating mice with monoamine oxidase type-A (MAO-A) and/or MAO-B inhibitors. Similar to MPTP, which is converted to MPP<sup>+</sup> by MAO-B, 2'-NH<sub>2</sub>-MPTP is hypothesized to be converted to a toxic pyridinium, 2'-NH<sub>2</sub>-MPP<sup>+</sup>. (b) (1) While 2'-NH<sub>2</sub>-MPTP appears to cross the blood-brain barrier, 2'-NH<sub>2</sub>-MPP<sup>+</sup> is not expected to do so based on findings with MPP<sup>+</sup>. 2'-NH<sub>2</sub>-MPTP toxicity is prevented by inhibiting plasma membrane serotonin transporters (SERT) or norepinephrine transporters (NET), which is hypothesized to block uptake of 2'-NH<sub>2</sub>-MPP<sup>+</sup> into serotonin and norepinephrine (NE) neurons, respectively. (3) Superoxide radicals are thought to be formed at mitochondria leading to membrane instability, DNA damage, and ultimately, axonal degeneration and to a lesser extent, cell death. Increasing superoxide dismutase activity prevents 2'-NH2-MPTP-induced toxicity. (4) Unlike MPP<sup>+</sup>, which is readily accumulated in synaptic vesicles via vesicular monoamine transporters type-2 (VMAT2), 2'-NH2-MPP+ appears to be a weaker substrate for VMAT2 and only low dose 2'-NH<sub>2</sub>-MPTP toxicity is potentiated by pharmacologic or genetic inhibition of VMAT2

2'-NH<sub>2</sub>-MPTP appeared warranted. However, the effects of 2'-NH<sub>2</sub>-MPTP on serotonin did not escape notice, and several years later, additional research established the unique neurotoxicity of 2'-NH<sub>2</sub>-MPTP (Andrews and Murphy 1993a, b, c).

# 3 2'-NH<sub>2</sub>-MPTP Is a Serotonin and Norepinephrine Neurotoxin in Mice

Initial investigation confirmed striatal serotonin loss in response to 2'-NH<sub>2</sub>-MPTP administration (Andrews and Murphy 1993a). Furthermore, 2'-NH<sub>2</sub>-MPTP was

shown to decrease norepinephrine (NE), in addition to serotonin in the forebrain, in contrast to the dopamine-selective toxicity produced by MPTP and its most potent analog, 2'-CH<sub>3</sub>-MPTP. When administered to C57Bl/6 mice, 2'-NH<sub>2</sub>-MPTP reduced serotonin, 5-HIAA, and norepinephrine levels by 70 % in frontal cortex and 60 % in hippocampus, with depletions occurring to a lesser extent in striatum and brain stem. Dopamine and its major metabolites, 3,4-dihydroxyphenylacetic acid (DOPAC) and homovanillic acid (HVA), were relatively unaffected 1 week after 2'-NH<sub>2</sub>-MPTP (Andrews and Murphy 1993a). Notably, no other MPTP analogs, including all amino-substituted and 2'-substituted analogs, have been reported to produce similar serotonin and norepinephrine neurotoxicity in mice.

#### 4 Acute Effects of 2'-NH<sub>2</sub>-MPTP

The median lethal dose  $(LD_{50})$  is the dose required to kill half of a treated population and is an indicator of the acute lethality of substances. The  $LD_{50}$  for 2'-NH<sub>2</sub>-MPTP in C57Bl/6 mice is 20–40 mg/kg, ip (Andrews and Murphy 1993a). It is likely to be similar in Swiss Webster mice (Andrews and Murphy 1993b) and somewhat lower in CD-1 mice (Andrews et al. 1996). When treated with high doses of 2'-NH<sub>2</sub>-MPTP, i.e.,  $\geq$ 40 mg/kg, mice had tonic-clonic seizures and died shortly afterwards (Andrews and Murphy 1993a). Tonic-clonic seizures followed by death in some cases also occurred in rats at 20 mg/kg 2'-NH<sub>2</sub>-MPTP (Unger et al. 2002). Acute and massive release of monoamine neurotransmitters has been shown to cause precipitous elevations in blood pressure and heart rate (Fuller et al. 1988). Death subsequent to high-dose 2'-NH<sub>2</sub>-MPTP administration is likely to be due to cardiovascular consequences and not neurodegeneration.

Similar to MPTP, the dose regimen most commonly used for 2'-NH<sub>2</sub>-MPTP in rodents consists of four intraperitoneal (ip) injections (15–20 mg/kg) at 2-h intervals. Behavioral changes in mice and rats immediately following each injection include piloerection, salivation, Straub (straight) tail, flattened body posture, intermittent walking, forepaw treading (piano-playing), and wet-dog shakes. Behavioral responses generally subside prior to each subsequent injection of 2'-NH<sub>2</sub>-MPTP and become less pronounced after repeated injections. Acute behavioral changes in rodents are characteristic of serotonin syndrome, which results from large increases in central serotonin release (Zhang et al. 2009). Serotonin syndrome arising from single drug overdose or multiple serotonergic agents acting via different but converging mechanisms is potentially life threatening in humans (Boyer and Shannon 2005).

Serotonin is also known to be involved in thermoregulation (Murphy et al. 1991). Acutely, 2'-NH<sub>2</sub>-MPTP causes a decrease in core body temperature in mice (Numis et al. 2004). This hypothermic effect is similar to that seen after MPTP (Freyaldenhoven et al. 1995; Moy et al. 1998) but opposite to substituted amphetamines, which cause hyperthermia (Miller and O'Callaghan 1995). It is not known why 2'-NH<sub>2</sub>-MPTP and substituted amphetamines, both of which are thought to cause acute serotonin release, have differing effects on core body temperature.

Loss of serotonin and norepinephrine in frontal cortex and hippocampus has been detected as early as 30 min after the last injection of 2'-NH<sub>2</sub>-MPTP in mice (Luellen et al. 2003). In contrast, decreases in 5-HIAA developed slowly over the first week after treatment. This probably reflects high levels of metabolism of serotonin from degenerating axons. In striatum, hypothalamus, and brain stem, where long-term depletions are more modest, serotonin and norepinephrine levels showed substantial recovery over the 72 h period following treatment. Transient decreases (30–40 %) in striatal dopamine were also reported during the initial 2 h after 2'-NH<sub>2</sub>-MPTP, but dopamine levels recovered by 24 h posttreatment. Thus, forebrain serotonergic and noradrenergic axons that project far from their origins in the brain stem show the most pronounced early evidence of toxicity. In addition to acute behavioral, physiological, and neurochemical changes, mice show a 1–2 g decrease in body weight during the first 48 h after 2'-NH<sub>2</sub>-MPTP. Throughout the first week after exposure, mice exhibit noise sensitivity and agitation.

## 5 Mechanism of Action of 2'-NH<sub>2</sub>-MPTP

While the mechanisms underlying 2'-NH<sub>2</sub>-MPTP-induced neurotoxicity are not fully understood, they have many similarities to those of MPTP. Both are small, lipophilic molecules that cross the blood-brain barrier (Fig. 1b). MPTP is a prodrug that requires oxidation by monoamine oxidase type-B (MAO-B) to exert toxicity as 1-methyl-4-phenylpyridinium (MPP<sup>+</sup>) (Markey et al. 1984; Heikkila et al. 1985). 2'-NH<sub>2</sub>-MPTP is oxidized by both MAO-B and monoamine oxidase type-A (MAO-A) (Fig. 1b) (Andrews and Murphy 1993b; Andrews 2000). Therefore, like MPTP, 2'-NH<sub>2</sub>-MPTP probably exerts its toxic effects through the oxidized pyridinium, 2'-amino-1-methyl-4-phenylpyridinium (2'-NH<sub>2</sub>-MPP<sup>+</sup>) (Fig. 1a). Preliminary findings show monoamine oxidase-dependent production of 2'-NH<sub>2</sub>-MPP<sup>+</sup> in brain tissue incubated with 2'-NH<sub>2</sub>-MPTP. 2'-NH<sub>2</sub>-MPP<sup>+</sup> was identified by tandem mass spectrometry and comparison with authentic 2'-NH<sub>2</sub>-MPP<sup>+</sup> via high-performance liquid chromatography.

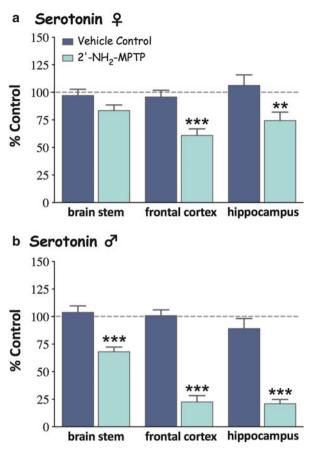
The MAO-catalyzed metabolite of 2'-NH<sub>2</sub>-MPTP gains access to serotonergic and noradrenergic neurons via serotonin transporters (SERT) and norepinephrine transporters (NET), respectively (Andrews and Murphy 1993b, c; Numis et al. 2004) (Fig. 1b), just as MPTP targets dopaminergic neurons via uptake of MPP<sup>+</sup> by dopamine transporters (DAT) (Melamed et al. 1985; Gainetdinov et al. 1997). Studies have shown that pretreating mice with the serotonin-selective reuptake inhibiting antidepressants fluoxetine or paroxetine prevent 2'-NH<sub>2</sub>-MPTP-induced serotonergic neurotoxicity (Andrews and Murphy 1993b, c). Pretreatment with the norepinephrine-selective reuptake inhibitor desipramine prevents norepinephrine depletion. 2'-NH<sub>2</sub>-MPTP neurotoxicity can also be limited by genetic inactivation of SERT to produce selective loss of norepinephrine (Numis et al. 2004). Administration of 2'-NH<sub>2</sub>-MPTP to mice genetically deficient in NET (Xu et al. 2000) has not been investigated.

Similar to MPTP, 2'-NH<sub>2</sub>-MPTP appears to cause neurotoxicity via generation of oxygen free radicals. MPP<sup>+</sup> has been shown to block the mitochondrial respiratory chain by inhibiting complex I (Cleeter et al. 1992) with two major consequences. First, conversion of NAD<sup>+</sup> to NADH is decreased, which interferes with mitochondrial proton gradients and subsequent production of ATP, depleting neurons of vital chemical energy. Second, metabolic disruption is purported to cause oxidative stress associated with an increase in oxyradicals and damage to DNA, proteins, and the cytoskeleton (Halliwell 1992). Mice genetically engineered to overexpress Cu-Zn superoxide dismutase, the soluble enzyme isoform that converts highly toxic superoxide radicals to hydrogen peroxide, were protected from 2'-NH<sub>2</sub>-MPTP toxicity, implicating superoxide mechanistically (Andrews et al. 1996). Together, findings on 2'-NH<sub>2</sub>-MPTP and MPTP suggest that inhibition of mitochondrial complex I by pyridiniums to generate oxyradicals is a primary mechanism of toxicity, not release and auto-oxidation of dopamine, which is specific to MPTP.

In addition to differential transporter specificity, there is another key difference between the molecular mechanisms of 2'-NH<sub>2</sub>-MPTP and MPTP. Previous studies have shown that MPP<sup>+</sup> is a substrate for the vesicular monoamine transporter type-2 (VMAT2), and sequestration of MPP<sup>+</sup> into synaptic vesicles reduces neurotoxicity (Gainetdinov et al. 1998; Staal and Sonsalla 2000b). Thus, MPP<sup>+</sup> mimics dopamine via high-affinity transport by DAT and VMAT2. In general, vesicular sequestration of exogenous toxins by VMAT2 has been implicated as one of several lines of neuronal defense (Miller et al. 1999). However, 2'-NH<sub>2</sub>-MPP<sup>+</sup> appears to be a poor substrate for VMAT2 (Numis et al. 2004). Mice with partial genetic reductions in VMAT2 expression, as well as mice treated with the VMAT2 inhibitor, Ro 4–1284 (2-hydroxy-2-ethyl-3-isobutyl-9,10-dimethoxy-1,2,3,4,6,7hexahydrobenzo[alpha]chinolizin hydrochloride), showed only slight potentiation of toxicity when treated with low doses of 2'-NH<sub>2</sub>-MPTP.

With regard to sex, female mice appear to be less susceptible to the neurotoxic effects of 2'-NH<sub>2</sub>-MPTP. Compared to male mice with 60–80 % loss of serotonin in frontal cortex and hippocampus 6 weeks after treatment, female CD-1 mice only showed 30–40 % serotonin depletions in the same brain regions (Fig. 2). Studies have shown that female mice are resistant to dopamine depletion after MPTP treatment (Miller et al. 1998; Dluzen 2000; Morissette et al. 2007). Castrated male mice pretreated with estrogen are also protected from MPTP toxicity (Dluzen et al. 1996). Estrogen may exert neuroprotective effects through direct antioxidant activity (Subbiah et al. 1993). However, these effects are only observed at low physiological concentrations of estrogen (Ramirez et al. 2003), implicating other mechanisms including gene regulation of estrogen receptors (Kenchappa et al. 2004) or altered DAT affinity for MPP<sup>+</sup> (Ookubo et al. 2009). Similar mechanisms may underlie sex-related differences in 2'-NH<sub>2</sub>-MPTP neurotoxicity.

Fig. 2 Changes in serotonin levels following 2'-NH2-MPTP treatment. Brain tissue neurotransmitter levels were analyzed by highperformance liquid chromatography with electrochemical detection. Data are percents of noninjected control levels in vehicle- and 2'-NH2-MPTPtreated mice. Serotonin levels were analyzed in (a) female mice and (b) male mice. Statistical significances: \*\*P < 0.01 and \*\*\*P < 0.0012'-NH2-MPTP-treated versus uninjected mice. N = 10 for control groups of both sexes; N = 9 for 2'-NH<sub>2</sub>-MPTPtreated group of both sexes



Thus, 2'-NH<sub>2</sub>-MPTP produces toxicity by cellular mechanisms with many similarities to those of MPTP, although different neuronal populations are targeted and the neuroprotective role of VMAT2 varies. Recently,  $\alpha$ -synuclein was found to be neuroprotective in the context of MPTP but not 2'-NH2-MPTP toxicity suggesting that  $\alpha$ -synuclein plays a role in dopaminergic toxicity but not serotonin and norepinephrine toxicity (Thomas et al. 2011). 2'-NH<sub>2</sub>-MPTP differs from other serotonin neurotoxins, such as 5,6- and 5,7-dihydroxytryptamine (5,6- and 5,7-DHT), which must be administered directly into the brain. 2'-NH<sub>2</sub>-MPTP is also mechanistically dissimilar from the norepinephrine neurotoxin DSP-4 (N-(2chloroethyl)-N-ethyl-2-bromobenzylamine hydrochloride) (Hallman and Jonsson 1984; Kostrzewa 2009) and the substituted amphetamines, e.g., 3,4-methylenedioxvlmethamphetamine, fenfluramine, and *p*-chloroamphetamine (Baumgarten and Lachenmayer 2004). The toxicity profiles of substituted amphetamines differ from 2'-NH<sub>2</sub>-MPTP, particularly in mice, where these amphetamines also affect dopamine levels (O'Callaghan and Miller 1994) (Jayanthi et al. 1999), or in the case of fenfluramine, where long-term neurotoxic effects are questionable.

## 6 Evidence for Neurodegeneration

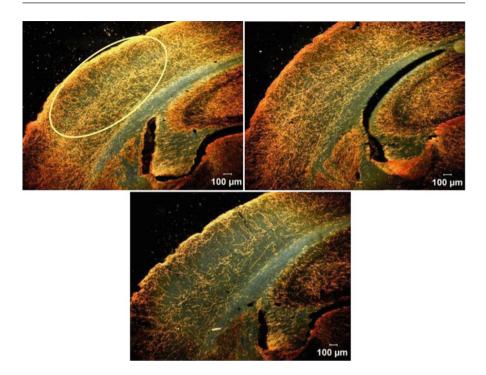
Multiple lines of investigation suggest that reductions in forebrain serotonin and norepinephrine are indicative of 2'-NH<sub>2</sub>-MPTP-induced neurodegeneration. This is compatible with findings of MPTP-induced dopaminergic neurodegeneration (Markey et al. 1986). Reductions in cortical and hippocampal serotonin and norepinephrine persist up to six months after treatment of mice with 2'-NH<sub>2</sub>-MPTP (Luellen et al. 2003). Furthermore, a number of indicators of neurodegeneration are present after 2'-NH<sub>2</sub>-MPTP administration.

When some types of neurons, including serotonin neurons, are damaged, i.e., during aging, disease, or injury, they increase production of brain-derived neurotrophic factor (BDNF), a protein involved in synaptic plasticity, neurogenesis, and neuronal survival (Mattson et al. 2004). Along these lines, 2'-NH<sub>2</sub>-MPTP-treated mice show acute increases in hippocampal BDNF (Szapacs et al. 2004; Luellen et al. 2006). Further evidence for degenerating neurons is hypertrophy of astrocytes in a process known as astrogliosis. Glial fibrillary acidic protein (GFAP) is an intermediate filament protein whose expression is universally upregulated in astrocytes in response to brain injury (O'Callaghan 1991). In addition to BDNF, 2'-NH<sub>2</sub>-MPTP administration to mice causes transient elevations in GFAP in frontal cortex and hippocampus, brain regions showing long-term serotonergic and noradrenergic depletion indicative of neuronal injury (Luellen et al. 2003). Interestingly, administration of 2'-NH<sub>2</sub>-MPTP to mice 2 weeks prior to treatment with the dopaminergic neurotoxin 2'-CH<sub>3</sub>-MPTP prevents striatal increases in GFAP in response to 2'-CH<sub>3</sub>-MPTP suggesting that serotonin is a modulator of astrogliosis (Luellen et al. 2003).

Immunocytochemistry has also been used to study neurodegeneration. A few days after 2'-NH<sub>2</sub>-MPTP treatment, serotonin-immunopositive axons exhibited a tortuous, beaded, and punctate appearance indicating ongoing neurodegeneration (Luellen et al. 2006). Three weeks later, significant serotonergic axonal losses were detected. The hippocampus and frontal cortex were the brain regions most severely affected, strongly correlating with reductions in neurotransmitter levels (Fig. 3). Silver staining has also been used for over a century as a histopathological indicator of neuronal death (Uchihara 2007). Immunocytochemical evidence of argyrophilia, the ability of degenerating neurons to take up silver, occurs in serotonin cell bodies in the dorsal raphe further indicating neurodegeneration after 2'-NH<sub>2</sub>-MPTP (Luellen et al. 2006). From these converging lines of evidence, it can be concluded that 2'-NH<sub>2</sub>-MPTP is a regionally selective, potent, and long-lasting serotonergic and noradrenergic neurotoxin.

## 7 2'-NH<sub>2</sub>-MPTP Toxicity in Rats and Nonhuman Primates

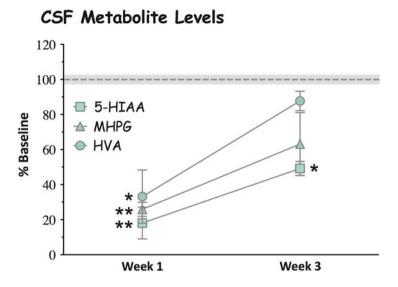
Notwithstanding the physiological and neurochemical similarities between 2'-NH<sub>2</sub>-MPTP and MPTP, different patterns of species sensitivity are observed. Notably, key differences between these two compounds occur regarding their effects in rats. Rats are virtually resistant to the effects of MPTP (Chiueh et al. 1984), a phenomenon that



**Fig. 3** Immunocytochemistry of serotonin axons following acute and chronic administration of 2'-NH<sub>2</sub>-MPTP. Representative dark field photomicrographs of sections of primary somatosensory cortex from male CD-1 mice are shown. Serotonin axons are gold threadlike structures. Mice were treated with  $4 \times 20$  mg/kg 2'-NH<sub>2</sub>-MPTP at 2 h intervals. Compared to control mice (*upper* left panel), mice sacrificed 21 days (*lower* panel) but not 3 days (*upper* right panel) post-2'-NH<sub>2</sub>-MPTP treatment showed significant reductions in serotonergic axon innervation in the barrel field region (*yellow* oval) (Reprinted with permission from (Luellen et al. 2006))

contributed to the difficulties of identifying MPTP as the cause of Parkinsonism in human drug users (Davis et al. 1979; Lewin 1984). However, systemic treatment of Sprague Dawley rats with 2'-NH<sub>2</sub>-MPTP ( $4 \times 15$  mg/kg, ip, every 2 h) was highly toxic, depleting forebrain serotonin and norepinephrine levels without affecting striatal dopamine (Unger et al. 2002). Conversely, 2'-CH<sub>3</sub>-MPTP-treated rats showed no toxic effects.

In terms of molecular mechanisms of MPTP and its analogs, there appear to be several ways in which species can differ. Some animals, rats in particular, have high MAO-B activity at the blood-brain barrier, where effective conversion of MPTP to MPP<sup>+</sup> can take place (Kalaria et al. 1987). Positively charged pyridinium species produced at the blood-brain barrier are then prevented from entering the brain (Fig. 1b). Rats may therefore be partly protected from MPTP toxicity by effective MPTP metabolism at the blood-brain barrier. Furthermore, rats have a greater capacity to sequester MPP<sup>+</sup> compared to mice (Staal and Sonsalla 2000a). However, 2'-NH<sub>2</sub>-MPP<sup>+</sup> appears to be a poor substrate for VMAT2 in

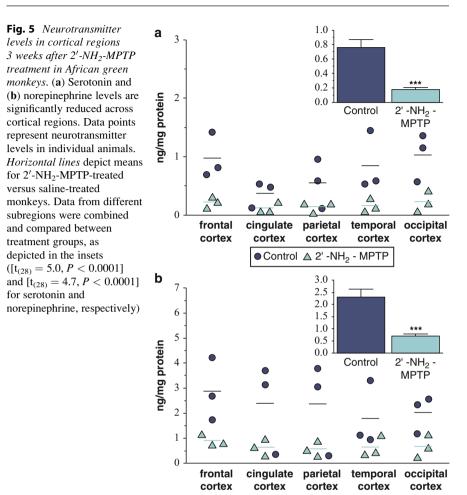


**Fig. 4** *Cerebrospinal fluid metabolite levels after administration of 2'-NH*<sub>2</sub>-*MPTP in African green monkeys.* Levels of 5-hydroxyindoleacetic acid (5-HIAA), 3-methoxy-4-hydroxyphenylglycol (MHPG), and homovanillic acid (HVA) were measured in cerebrospinal fluid (CSF) 1 and 3 weeks after administration of 2'-NH<sub>2</sub>-MPTP to African *green* monkeys (20 mg/kg total dose). These are metabolites of serotonin, norepinephrine, and dopamine, respectively. All metabolites were decreased 1 week after treatment with 2'-NH<sub>2</sub>-MPTP and remained reduced for 5-HIAA 3 weeks after treatment. The *gray-shaded* region represents  $\pm$  the standard error of the mean (SEM) for all metabolites in saline-treated monkeys. Data are means  $\pm$  SEMs. \**P* < 0.05 and \*\**P* < 0.01 versus metabolite levels in saline-treated animals

mice (Numis et al. 2004) and rats (Unger et al. 2002). Lower blood–brain barrier metabolism of 2'-NH<sub>2</sub>-MPTP and synaptic vesicular uptake of 2'-NH<sub>2</sub>-MPP<sup>+</sup> are hypothesized to contribute to the sensitivity of rats to 2'-NH<sub>2</sub>-MPTP toxicity.

In addition to rodents, 2-NH<sub>2</sub>-MPTP has been administered to two nonhumanprimate species. In one study, six adult African green monkeys (*C. aethiops*) were given intravenous (iv) 2'-NH<sub>2</sub>-MPTP with the objective of investigating serotonin and norepinephrine neurotoxicity in a species more closely related to humans. Initial doses of 2'-NH<sub>2</sub>-MPTP (7–10 mg/kg) to one monkey produced behavior indicative of hallucinations followed by nonlethal tonic-clonic seizure at the highest dose. A subsequent dose of 5 mg/kg to the same animal was without overt behavioral effects. Two additional monkeys received  $4 \times 5$  mg/kg 2'-NH<sub>2</sub>-MPTP. Bolus iv doses were administered at a 6 h interval twice per day for two consecutive days. This dosing schedule of 2'-NH<sub>2</sub>-MPTP is similar to other serotonin neurotoxin regimens in nonhuman primates (Ricaurte et al. 1988).

Cerebrospinal fluid (CSF) was analyzed after 2'-NH<sub>2</sub>-MPTP to assess changes in neurotransmitter metabolite levels (Fig. 4). The serotonin and norepinephrine metabolites 5-HIAA and 3-methoxy-4-hydroxyphenylglycol (MHPG), respectively, were significantly reduced 1 week after 2'-NH<sub>2</sub>-MPTP treatment compared to levels in



saline-treated monkeys. Depressed CSF 5-HIAA levels continued to be evident at 3 weeks posttreatment with 2'-NH<sub>2</sub>-MPTP. The dopamine metabolite HVA was also moderately reduced 1 week after drug treatment. Changes in CSF metabolite levels could indicate loss of brain serotonergic and noradrenergic (and possibly dopaminergic) axons/neurons. They might also reflect inhibition of MAO activity. Studies have shown that when MPTP is oxidized to form MPP<sup>+</sup>, MAO-B is irreversibly inactivated (Singer et al. 1986). "Suicide" inhibition of MAO is expected to reduce monoamine neurotransmitter metabolites. Notably, MAO inhibition by 2'-NH<sub>2</sub>-MPTP has not yet been investigated. In addition to CSF metabolites, serotonin and norepinephrine levels across cortical regions were globally decreased in African green monkeys treated 3 weeks earlier with 2'-NH<sub>2</sub>-MPTP (Fig. 5) suggesting that like MPTP, nonhuman primates are susceptible to 2'-NH<sub>2</sub>-MPTP neurotoxicity. In African green monkeys, brain dopamine levels were modestly affected in the caudate (-24 %) but not the putamen (-9 %).

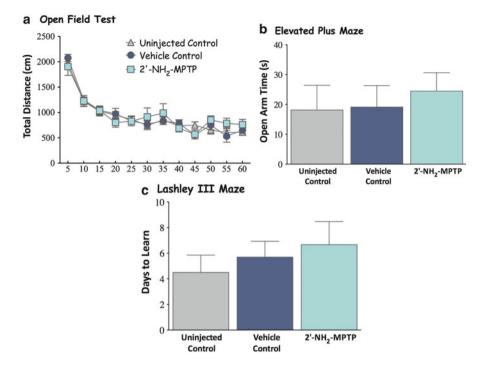
2'-NH<sub>2</sub>-MPTP has also been administered to four adult baboons (*P. anubis*). Here, the neurotoxin was infused unilaterally into the carotid artery. This route of administration has been used with MPTP resulting in dopaminergic neurodegeneration largely confined to the ipsilateral side of the brain (Bankiewicz et al. 1986; Guttman et al. 1990). Intracarotid administration of MPTP produces a hemiparkinsonian nonhuman primate model where animals can still feed and care for themselves (Ding et al. 2008). Additionally, one hemisphere is relatively unaffected and serves as an intra-animal control in morphological studies (Przedborski et al. 1991). Intracarotid administration of 2'-NH<sub>2</sub>-MPTP to baboons was carried out to attempt to produce a model of unilateral loss of forebrain serotonergic and noradrenergic axons that could be used to evaluate positron emission tomography (PET) ligands. Like unilateral MPTP, each 2'-NH<sub>2</sub>-MPTP-treated animal would serve as its own control in imaging studies.

A 10 mg/kg infusion of 2'-NH<sub>2</sub>-MPTP to a single baboon resulted in nonlethal seizure. Doses of 2'-NH<sub>2</sub>-MPTP in the remaining baboons were lowered to 6–7 mg/kg. Bilateral biopsies of cortex suggested unilateral reductions in serotonin and norepinephrine. However, the data were inconclusive due to large variations in cortical neurotransmitter levels across samples. Postmortem brain tissue analysis showed evidence for loss of dopamine in the caudate and putamen of 2'-NH<sub>2</sub>-MPTP-treated baboons. One baboon treated unilaterally with 2'-NH<sub>2</sub>-MPTP underwent PET scanning using [<sup>11</sup>C]-McN5652 to label SERT in extant neurons (Parsey et al. 2000). SERT density was unilaterally reduced across all brain regions except in the cerebellum. While some studies find SERT (albeit at extremely low levels), and 5-HT<sub>1A</sub>, 5-HT<sub>2A</sub>, and 5-HT<sub>3A</sub> receptors in the cerebellum, direct evidence that 5-HT neurons innervate this region remains inconclusive (Kish et al. 2005; Arpin-Bott et al. 2006; Xie et al. 2006; Yew et al. 2009). In fact, many investigators use the cerebellum to define nonspecific binding when analyzing SERT levels.

Together, findings in nonhuman primates suggest that similar to rats and mice, 2'-NH<sub>2</sub>-MPTP may be neurotoxic to serotonin and norepinephrine neurons in primate species. These data also suggest minor dopaminergic neurotoxicity of 2'-NH<sub>2</sub>-MPTP in nonhuman primates. The latter might be due to the routes of administration or high doses of 2'-NH<sub>2</sub>-MPTP used in monkeys. Small groups of animals and inherent variability across individuals also contributed to the incomplete yet highly suggestive nature of these first studies on 2'-NH<sub>2</sub>-MPTP in primates.

## 8 Long-Term Behavioral Consequences of 2'-NH<sub>2</sub>-MPTP

Compared to research on the neurotoxic mechanisms of 2'-NH<sub>2</sub>-MPTP in different animal models, relatively little work has been done to understand the behavioral consequences of serotonergic and/or noradrenergic forebrain axonal loss. The effects of 2'-NH<sub>2</sub>-MPTP neurotoxicity were recently studied on locomotor activity (Unger et al. 2006), anxiety-related behavior (Walf and Frye 2007), and low-stress spatial learning and memory (Bressler et al. 2010). Here, CD-1 mice were pretreated with desipramine to produce serotonin-selective



**Fig. 6** *Effects of* 2'-*NH*<sub>2</sub>-*MPTP treatment on behavior in male mice.* (a) Locomotor behavior was assessed in open field activity monitors (40 cm × 40 cm × 30 cm), where mice were able to explore freely for 1 h. No changes in total distance traveled were associated with 2'-NH<sub>2</sub>-MPTP treatment. (b) The elevated plus maze was used to investigate changes in anxiety-related behaviors as indicated by time spent in the aversive open arms of the maze. No changes in anxiety-related behavior were observed after 2'-NH<sub>2</sub>-MPTP. (c) Low-stress route learning was investigated via the Lashley III maze. A mouse was considered as having learned the maze (reached criterion) after completing two consecutive nightly trials with only 0 errors or 1 error. Mice were tested once per night for seven consecutive nights. Mice treated with 2'-NH<sub>2</sub>-MPTP (N = 9) took a similar number of days to learn the maze compared with unhandled (N = 10) and vehicle controls (N = 10) indicating intact learning capabilities

2'-NH<sub>2</sub>-MPTP-induced toxicity. It was hypothesized that serotonergic axonal degeneration would be associated with impaired cognition and increased anxiety-related behavior, similar to changes in brain function observed in humans with neurodegenerative disorders having serotonin-system involvement. Nonetheless, 60–70 % reductions in serotonin in frontal cortex and hippocampus (Fig. 2) were not associated with changes in locomotor, anxiety-related, and learning behavior in the open field, elevated plus maze, or Lashley III maze, respectively (Fig. 6).

Studies have shown that disruption of the serotonin system in response to another serotonin neurotoxin, 5,7-DHT, led to changes in learning and memory in rodents and nonhuman primates (Lieben et al. 2006; Clarke et al. 2007), although not all studies have been in agreement (Adams et al. 2008). Anxiety-like behaviors in 5,7-DHT-treated animals were not altered (Lieben et al. 2006). Here, adult mice ages 12–16 weeks were studied. Thus, critical early or late windows of vulnerability could

have been missed when degeneration of forebrain serotonin axons influences behavior. Alternately, behavioral changes might occur in 2'-NH<sub>2</sub>-MPTP-treated mice, but the tests focused on were not directed at putative behavioral changes, e.g., depression-related behavior. Further research will also need to be carried out with 2'-NH<sub>2</sub>-MPTP in genetically engineered disease-specific models to understand the ramifications of serotonin and norepinephrine degeneration on behavior.

#### 9 Conclusion

A number of studies show that 2'-NH2-MPTP induces significant serotonergic and noradrenergic neurotoxicity in mice, rats, and probably, nonhuman primates. The mechanisms of 2'-NH<sub>2</sub>-MPTP neurotoxicity are similar to those of MPTP, yet significant differences occur with regard to the isoforms of monoamine oxidase that produce proximal pyridinium neurotoxins; 2'-NH<sub>2</sub>-MPTP is a substrate for MAO-A and MAO-B while MPTP is metabolized almost exclusively by MAO-B. Moreover, rats are relatively insensitive to the dopamine-depleting effects of MPTP (and 2'-CH<sub>3</sub>-MPTP). Conversely, 2'-NH<sub>2</sub>-MPTP produces serotonin and norepinephrine toxicity in rats at doses comparable to those used in mice. The latter is likely to be associated, at least in part, to differential affinities of MPP<sup>+</sup> versus 2'-NH<sub>2</sub>-MPP<sup>+</sup> for VMAT2 in rats and mice. The most important differences between 2'-NH<sub>2</sub>-MPTP and MPTP arise from their affinities for plasma monoamine uptake transporters. This mechanistic aspect bestows relative dopamine-selective neurotoxicity to MPTP and 2'-CH<sub>3</sub>-MPTP. By contrast, the MAO-catalyzed oxidative metabolite of 2'-NH<sub>2</sub>-MPTP is a substrate for serotonin and norepinephrine transporters, conferring 2'-NH<sub>2</sub>-MPTP with the ability to produce serotonergic and noradrenergic neurotoxicity largely in the absence of dopaminergic neurotoxicity. Further research in nonhuman primates and with regard to the behavioral consequences of serotonergic and noradrenergic neurodegeneration seems to be warranted. In turn, 2'-NH<sub>2</sub>-MPTP animal models could be used to investigate potential treatments for serotonin-related pathologies. Moreover, a clearer understanding of links between serotonin deficiency and behavior may ultimately help to unravel the complex pathophysiology of depression and other neuropsychiatric and neurodegenerative disorders.

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# Methamphetamine and MDMA Neurotoxicity: Biochemical and Molecular Mechanisms

# Verónica Bisagno and Jean Lud Cadet

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#### Abstract

Methamphetamine (METH) and its analogs, methylenedioxymethamphetamine (MDMA), are psychostimulant drugs with high abuse liability. The two drugs are also very neurotoxic. In the case of METH, the behavioral and neurotoxic effects of the drug occur because it alters dopamine terminal physiology and causes massive release of dopamine (DA) in the synaptic cleft in brain regions that receive dopaminergic projections from

V. Bisagno

J.L. Cadet (🖂) NIDA Intramural Program, Molecular Neuropsychiatry Research Branch, NIDA/NIH/DHHS, Baltimore, MD, USA e-mail: jcadet@intra.nida.nih.gov

Instituto de Investigaciones Farmacológicas (ININFA-UBA-CONICET), Buenos Aires, Argentina e-mail: vbisagno@ffyb.uba.ar

the midbrain. The increase of synaptic DA is compounded by the ability of METH to block DA reuptake into DA terminals. METH toxicity is not only accompanied by terminal dysfunction but also by causing dysfunction of complex networks that subserve cognitive and emotional processes. MDMA is a ring-substituted derivative of phenylisopropylamine which is structurally similar to METH. MDMA is a substrate of the serotonin transporter (SERT) via which it enters monoaminergic neurons and causes release of serotonin (5-HT) from storage vesicles. This is followed by 5-HT release into the synaptic cleft by reversal of normal SERT function. MDMA is selectively neurotoxic to serotonergic nerve terminals in rats, guinea pigs, and nonhuman primates. MDMA users consistently show reduced SERT radionuclide ligand binding across multiple brain regions. There is also evidence that MDMA users can suffer from cognitive deficits. However, the relation of DA and/or 5-HT depletion to cognitive impairments remains to be clarified in METH and MDMA users. Results from these studies are likely to impact the therapeutic approaches to the treatment of patients who suffer from METH and MDMA addiction.

List of Abbre	viations
5-HIAA	5-hydroxyindoleacetic acid
5-HT	Serotonin
α-MT	α-methyl-p-tyrosine
CNS	Central nervous system
COMT	Catechol-O-methyltransferase
CuZnSOD	Copper-zinc superoxide dismutase
DA	Dopamine
DAT	Dopamine transporter
ER-	Endoplasmic reticulum-
HHA	3,4-Dihydroxyamphetamine
HHMA	3,4-Dihydroxymethamphetamine
HMA	4-Hydroxy-3-methoxy-amphetamine
L-DOPA	L-3,4-Dihydroxyphenylalanine
MDA	Methylenedioxyamphetamine
MDMA	Methylenedioxymethamphetamine
METH	Methamphetamine
MK-801	Dizocilpine
MTF-1	Metal-responsive transcription factor 1
MTs	Metallothioneins
NET	Norepinephrine transporter
Nrf2	NF-E2-Related factor 2
PARP	Poly(ADP-ribose) polymerase
ROS	Reactive oxygen species
SERT	Serotonin transporter
TH	Tyrosine hydroxylase
TPH	Tryptophan hydroxylase
VMAT2	Vesicular monoamine transporter 2

## 1 Introduction

Some psychostimulants including methamphetamine (METH) are presently prescribed for weight control, narcolepsy, and attention deficit disorder. These drugs are also used recreationally, with some individuals developing abuse followed by severe addiction. METH addiction is prevalent throughout the world, being second only to marijuana abuse among illicit drugs. During the past two decades, it has become clear that dopamine (DA) plays a critical role in the behavioral pharmacology and addictive properties of abused drugs. DA contributes to incentive salience and is necessary for identifying wanted elements from the environment (Berridge 2007). Drugs of abuse that increase dopaminergic signals alter and sensitize dynamic mesolimbic mechanisms that evolved to attribute incentive salience to rewards (Le Moal and Koob 2007). Such drugs interact with incentive salience integrating Pavlovian associative information with physiological state signals (Berridge 2007). Specific transporters in the synaptic membrane of dopaminergic terminals serve to remove DA from the synaptic cleft to end its action (Sulzer et al. 2005). METH alters dopamine terminal physiology and causes massive release of DA in the synaptic cleft in brain regions that receive dopaminergic projections from the midbrain (Krasnova and Cadet 2009). This is compounded by its ability to block DA reuptake into DA terminals. These mechanisms work together to cause degeneration of monoaminergic terminals in the striatum (O'Callaghan and Miller 1994; Krasnova and Cadet 2009). In the following paragraphs, we describe specific toxic effects and the mechanisms by which METH and its analogs, methylenedioxymethamphetamine (MDMA), induce damage of monoaminergic terminals and non-aminergic cell bodies in the brain.

# 2 Methamphetamine Toxicity

## 2.1 Dopamine and Oxidative Stress

Methamphetamine enters DA neurons through the DAT and via passive diffusion. In the cell, it accumulates in vesicles, disrupts the pH gradient required for vesicular DA sequestration, and displaces DA into the cytoplasm (Sulzer and Rayport 1990). METH causes release of monoamines from the neuronal cytosol via plasmalemmal uptake transporters, particularly the DA transporter (DAT), the norepinephrine transporter (NET), and the serotonin (5-HT) transporter (SERT) via a mechanism referred to as reverse transport (Sulzer et al. 2005). METH acts through the vesicular monoamine transporter 2 (VMAT-2) to cause excessive release of dopamine into the cytoplasm followed by DA release into the synaptic cleft through the DAT (Sulzer et al. 2005).

DA accumulates in the cytoplasm which alters the concentration gradient and likely helps to favor the reverse transport of DA via the DAT (see Sulzer et al. 2005).

DA is an important component of the mechanisms that underlie METH neurotoxicity (Sulzer and Rayport 1990).

Neurotoxic effects of METH have been demonstrated in many species including rats, mice, guinea pigs, cats, and monkeys (for detailed reviews see McCann and Ricaurte 2004 and Krasnova and Cadet 2009). Animals given repeated doses of METH suffer significant loss of DA and 5-HT in the striatum, cortex, and olfactory bulb (Deng et al. 2007) and decreases in striatal DAT binding, DAT immunoreactivity and tyrosine hydroxylase (TH) immunoreactivity (Bowyer et al. 2008; Deng et al. 1999), and TH protein levels (O'Callaghan and Miller 1994). Animals that self-administer METH also show deficits in dopaminergic systems (Krasnova et al. 2010). Morphological studies suggest that the reductions in the markers of DA and 5-HT system integrity are related to degeneration of DA and 5-HT axonal terminals (Ricaurte et al. 1982). These neurotoxic consequences can be prevented or attenuated by agents, such as  $\alpha$ -methyl-p-tyrosine, that block DA synthesis and decrease DA levels in the striatum (Kuhn et al. 2008). In contrast, treatment with immediate DA precursor, L-3,4-dihydroxyphenylalanine (L-DOPA), that restores cytoplasmic DA levels enhances METH toxicity (Kuhn et al. 2008).

Autoxidation and metabolism of synaptic DA can lead to generation of reactive oxygen species (ROS) including hydrogen peroxide and superoxide radicals, quinones, and semiquinones (Cadet 1988; Cadet and Brannock 1998). The involvement of superoxide radicals in METH toxicity on DA terminals has been tested in transgenic mice that overexpress the human copper–zinc superoxide dismutase (CuZnSOD), a cytosolic enzyme that catalyzes the breakdown of super-oxide radicals (Cadet et al. 1994a, b; Jayanthi et al. 1998). These mice that have much higher CuZnSOD enzyme activity in the cytosol than control wild-type animals (Jayanthi et al. 1998) are protected against METH toxicity (Cadet et al. 1994a, b).

METH can also cause neuronal apoptosis in several brain regions, including the striatum, cortex, hippocampus, and olfactory bulb (Deng and Cadet 2000; Deng et al. 2001, 2007; Ladenheim et al. 2000). Because METH administration is associated with ROS production and because ROS can cause apoptosis and DNA damage (Li and Trush 1993), we thought it is likely that METH might induce neuronal apoptosis through ROS-mediated DNA damage. This idea was supported by the observations that METH-induced poly(ADP-ribose) polymerase (PARP) cleavage, increase in caspase-3 activity, and neuronal death were all attenuated in the striata of CuZnSOD transgenic mice (Deng and Cadet 2000). A role for oxidative stress in METH toxicity is supported by data obtained from microarray analyses that have identified changes in the expression of a number of genes that participate in DNA repair, including APEX, PolB, LIG1, and DNA mismatch repair proteins MSH3 and PMS1 after toxic doses of the drug (Cadet et al. 2002).

METH treatment causes alterations in the DAT, including the formation of higher molecular mass DAT-associated complexes (Hadlock et al. 2009). This phenomenon is attenuated by either prevention of METH-induced hyperthermia

or pretreatment with the DA synthesis inhibitor,  $\alpha$ -methyl-p-tyrosine ( $\alpha$ -MT) (Baucum et al. 2004). Reactive species also are implicated in DAT complex formation because in vitro exposure to the reducing agent,  $\beta$ -mercaptoethanol, reverses this process (Baucum et al. 2004). Concurrent with METH-induced DAT complex formation, METH treatment causes a loss of DAT monomer immuno-reactivity and a decrease in DAT function (Baucum et al. 2004).

Another aspect of METH toxicity is the fact that the drug can cause transcription factor NF-E2-related factor 2 (Nrf2) transit cytosolic into nuclear fractions where the protein exerts its regulatory functions (Jayanthi et al. 2009). Nrf2 and metal-responsive transcription factor 1 (MTF-1) are known to play a key role in transcriptional activation of the MT-1/2 genes in response to heavy metals or oxidative stress (Miyazaki et al. 2010), and it has been suggested that astrocytes can act to protect surrounding neurons against excess DA and DA-quinone formation by upregulating quinone-quenching molecules like metallothioneins (MTs) (Miyazaki et al. 2010). Increases in DA might promote nuclear translocation of Nrf2, upregulating MT1/2 expression in astrocytes, suggesting a protective role of Nrf2 against METH toxicity. Accordingly, Granado et al. (2011b) showed that Nrf2 deficiency exacerbated METH-induced damage to dopamine neurons (increase in loss of TH and DAT-containing fibers in striatum) and also potentiated glial activation.

## 2.2 Role of DAT and VMAT-2 in METH Toxicity

The role of DAT in METH toxicity has been investigated using both pharmacological and genetic means. For example, administration of the DAT inhibitor, methylphenidate, 1 hour after METH treatment, can reverse decreases in vesicular DA uptake, reductions in VMAT-2 ligand binding, and decreases in VMAT-2 immunoreactivity in vesicular subcellular fractions 6 h after injections of the drugs (Sandoval et al. 2003). Modafinil, an anti-narcoleptic drug that has been used off-label to treat psychostimulant addiction, acts as a DAT blocker (Zolkowska et al. 2009) and is neuroprotective against METH toxicity, as documented by attenuation of METH-induced decreases in TH immunoreactivity and DA striatal content (Raineri et al. 2011). Importantly, mice with genetic deletion of DAT are also protected against drug-induced DA depletion, reactive astrocytosis, and ROS production in the striatum (Fumagalli et al. 1998).

A role for VMAT-2 in METH-induced damage to striatal DA terminals is also supported by studies showing that pretreatment with the irreversible inhibitor of vesicular transport, reserpine exacerbates toxicity of the psychostimulant (Kuhn et al. 2008). Additionally, the administration of the alkaloid, lobeline, after injections of METH was able to reverse METH-induced decreases in synaptosomal, membrane-associated, and vesicular VMAT-2 immunoreactivity 24 h after drug treatment while also preventing METH-induced striatal depletion (Eyerman and Yamamoto 2005).

### 2.3 Role of Dopamine Receptors in METH Toxicity

Several studies have also documented a role for DA receptors in the mediation of METH toxicity (Angulo et al. 2004; Beauvais et al. 2011; Javanthi et al. 2005; Xu et al. 2005). Specifically, pretreatment with the DA  $D_1$  receptor antagonist, SCH23390 [R-(+)-7-chloro-8-hydroxy-3-methyl-1-phenyl-2,3,4,5tetrahydro-1H-3-benzazepine], given before each of 5 METH injections was able to attenuate drug-induced decreases in TH activity and DA levels in the rat striatum and decreases in tryptophan hydroxylase (TPH) activity and 5-HT levels in the striatum and cortex measured 18-20 h after METH treatment (Sonsalla et al. 1986). The DA D<sub>2</sub> receptor antagonist, sulpiride, also blocked METH-induced toxic effects on DA systems without affording any protection to striatal and cortical 5-HT terminals (Sonsalla et al. 1986). In addition, a single injection of the DA D<sub>1</sub> receptor antagonist, SCH23390, prior to a single highdose METH injection also attenuated long-term decreases in DA levels (Jayanthi et al. 2005), reductions in DAT binding, and depletion in TH protein levels and reactive astrocytosis (Xu et al. 2005). SCH23390 also protected against METH-induced cell death in the striatum (Jayanthi et al. 2005; Xu et al. 2005). The protection afforded by SCH23390 against METH-mediated cell death appears to depend, in part, on the activation of the Fas/FasL death pathway because pretreatment with SCH23390 caused suppression of METHinduced increases in the expression of FasL and caspase-3 in rat striatal cells (Jayanthi et al. 2005). Jayanthi et al. (2009) also measured the expression of several genes that are regulated during ER stress. These include ATF3, HSP27, Hmox1, HSP40, CHOP/Gadd153 and ATF4, ATF6, BiP/GRP78, and GADD34. The DA  $D_1$  receptor antagonist also attenuated or blocked METH-induced increases in the expression of the majority of these genes. METH injections can also cause substantial increases in the expression of proteins that participate in endoplasmic reticulum- (ER-) and mitochondrial-dependent stress responses, and these changes were also blocked by SCH23390 (Beauvais et al. 2011). However, the D<sub>2</sub>-like receptor antagonist, raclopride, was found to have only small to moderate effects on ER stress proteins and only small effects on mitochondrial-dependent cellular stress proteins (Beauvais et al. 2011). Both DA  $D_1$  and  $D_2$  receptors appear to be required for METH toxicity since both D<sub>1</sub>R (Ares-Santos et al. 2012) and D<sub>2</sub>R (Granado et al. 2011a) knockout mice are protected against METH neurotoxicity.

#### 2.4 Role of Microglia on METH-Induced Toxicity

METH toxicity is accompanied by reactive astrocytosis (Bowyer et al. 1994) and microglial activation (Pubill et al. 2002) in various brain regions. In the normal brain, microglial cells exist in a resting state, but, in response to inflammation or brain damage, these cells increase in size, migrate to the site of the injury, and phagocytize dying and dead cells (Raivich 2005). Although microglial activation is

important for immune responses, their overactivation can result in neurotoxic consequences. This process seems to be involved in the case of METH toxicity. For example, METH induces microgliosis in areas of the brain that suffer from neuronal degeneration (Thomas and Kuhn 2005a; Thomas et al. 2004a, b). Reserpine and clorgyline that exacerbate METH toxicity also potentiate in METH-induced microglial activation in the mouse striatum (Thomas et al. 2008). In contrast, attenuation of METH neurotoxicity by dizocilpine (MK-801), dextromethorphan, and  $\alpha$ -methyl-p-tyrosine results in inhibition of METH-induced microglial activation (Kuhn et al. 2008; Thomas and Kuhn 2005b). Anti-inflammatory drugs, ketoprofen and indomethacin, and the second-generation tetracycline minocycline afford protection against METH-induced toxicity and microgliosis (Asanuma et al. 2003, 2004; Zhang et al. 2006). Further studies are necessary to identify the specific mechanisms that underlie METH-induced microgliosis and the manner by which these activated microglial cells might cause degeneration of monoaminergic terminals.

## 3 Preconditioning Doses of METH Protect Against Toxic Doses of METH

Pretreatment with multiple low-dose injections of METH or with gradually escalating doses of the drug has been shown to provide partial protection against the deleterious effects of a high-dose METH challenge (Cadet et al. 2009, 2011; Danaceau et al. 2007; Graham et al. 2008). These investigators use a regimen that involves gradual increases in METH administration to rats in order to mimic progressively larger doses of the drug used by some human METH addicts. METH preconditioning was associated with complete protection against dopamine depletion caused by a METH challenge in the striatum and cortex (Cadet et al. 2009). Hodges et al. (2011) also reported that this regimen resulted in prolonged protection, with a second challenge of METH causing no further decreases in striatal DA or 5-HT levels in comparison to the single METH challenge. Another study has also reported that, although it was associated with some decreases in striatal DAT, METH self-administration attenuated the persistent deficits in dopaminergic neuronal function and increases in GFAP immunoreactivity caused by a subsequent binge METH exposure (McFadden et al. 2012).

These data suggest that repeated injections of low doses of METH might render the brain refractory to the toxic effects of binge injections of the drug, thus providing protection against toxic METH effects in various brain regions. Neuroprotection induced by METH preconditioning might be related to reductions in METH hyperthermia (McFadden et al 2012). Another possible explanation for the neuroprotective effects of METH is the fact that, by reducing striatal DAT and increasing high-molecular DAT complexes that are nonfunctional, the striatum might have become less susceptible to the toxic effects of a METH challenge (Krasnova et al. 2011). This line of reasoning is consistent with the recent report that a low, sub-toxic dose of METH was able to attenuate 6-OHDA-induced decreases in ATP levels and mitochondrial dehydrogenase activity (El Ayadi and Zigmond 2011).

## 4 Summary of Discussion of METH Toxicity

Chronic abuse of METH contributes to depression, psychosis, mood disturbances, and psychomotor dysfunctions in humans (Darke et al. 2008; Homer et al. 2008). METH addicts also suffer from deficits in attention, working memory, and decision-making (Gonzalez et al. 2004; Verdejo-García et al. 2006). Withdrawal from the drug is also associated with anhedonia, irritability, and intense craving for the drug (Darke et al. 2008; Homer et al. 2008). Human addicts also suffer persistent decreases in the levels of DAT in the orbitofrontal cortex, dorsolateral prefrontal cortex, and the caudate-putamen (McCann et al. 1998; Sekine et al. 2003; Volkow et al. 2001a, c). METH abusers also show abnormal glucose metabolism in cortical and subcortical brain areas (Volkow et al. 2001b; Wang et al. 2004). In addition, there is prominent microglial activation in the brains of these individuals (Sekine et al. 2008). Alterations in DA tone produced by METH toxicity on dopamine terminals might not only influence the physiology of the brain but might also alter the complex networks that subserve cognitive and emotional processes. It is thus likely that some of the clinical observations in METH abusers might be secondary to METH-induced toxic and neuroinflammatory responses, suggesting that the development of therapeutic agents that address these processes might be necessary to address the basic neurobiology of METH-induced neuropsychiatric manifestations.

## 5 MDMA Neurotoxicity

MDMA is a ring-substituted derivative of phenylisopropylamine, structurally similar to METH and the hallucinogen, mescaline (Lyles and Cadet 2003). MDMA impacts peripheral and central nervous system (CNS) functions through its actions on the serotonergic system (Baumann and Rothman 2009). MDMA is a substrate of the serotonin transporter via which it enters monoaminergic neurons and causes release of 5-HT from storage vesicles, followed by 5-HT release into the synaptic cleft by reversal of normal SERT function (Baumann and Rothman 2009). Injections of large doses of MDMA cause massive release of 5-HT from presynaptic vesicles, followed by a rapid decrease in 5-HT and 5-hydroxyindoleacetic acid (5-HIAA) levels and decreased TPH activity (Lyles and Cadet 2003). There do not appear to be losses of 5-HT uptake sites at early time points after MDMA administration (Lyles and Cadet 2003). MDMA can also cause dose-dependent DA and NE release in the striatum and hippocampus in rats (Lyles and Cadet 2003). Adverse effects of acute MDMA administration, including cardiovascular stimulation and elevated

body temperature, are thought to involve monoamine release from sympathetic nerves in the periphery or nerve terminals in the CNS (Baumann and Rothman 2009). MDMA is selectively neurotoxic to serotonergic nerve terminals in rats, guinea pigs, and nonhuman primates (Green et al. 1995). In mice, however, MDMA appears to affect mainly the nigrostriatal dopaminergic system (Cadet et al. 1995). MDMA users consistently show reduced SERT radionuclide ligand binding across multiple brain regions (Reneman et al. 2006). Some evidence has suggested that these levels began to recover with increased periods of MDMA abstinence. Recently, Erritzoe et al (2011) reported significantly lower SERT binding potential in the neocortex (-56 %), pallidostriatum (-19 %), and amygdala (-32 %), with the extent of binding correlated with lifetime MDMA usage. Kish et al (2010) compared moderate Ecstasy/MDMA users with nonuser controls and reported that SERT binding was significantly reduced in all the cerebral cortices and hippocampus. The degree of these reductions was significantly associated with the extent of past MDMA usage. An additional molecular structural component of the serotonergic signaling pathway is the  $5-HT_{2A}$ receptor. 5-HT<sub>2A</sub> receptors are densely distributed in human cortex; 5-HT<sub>2A</sub> receptor levels were reduced in multiple brain regions in the ongoing/acute MDMA user group, possibly because of ongoing MDMA-mediated increases in 5-HT release (Reneman et al. 2002).

Neurochemical and anatomical studies have reported long-term reductions in markers of 5-HT systems in rats. These include decreased levels of 5-HT and of its major metabolite, 5-HIAA; decreased number of 5-HT transporters; and decreased activity of the rate-limiting enzyme of 5-HT synthesis, TPH (Commins et al. 1987; De Souza et al. 1990). Severe reductions in 5-HT and 5-HIAA levels occurred in the rat neocortex, striatum, and hippocampus (De Souza et al. 1990). These abnormalities are reported to last for months or even years after drug administration (Lyles and Cadet 2003). MDMA also perturbs the function of SERT (Green et al. 2003), a marker of the integrity of serotonin neurons (Blakely et al. 1994). By virtue of its moderating synaptic 5-HT levels, SERT is crucial for the process of 5-HT neurotransmission (Green et al. 2003). MDMA downregulates SERT function without altering SERT mRNA or protein expression, and this rapid downregulation is sustained for at least 90 min and is dose dependent (Kivell et al. 2010).

Histologic studies in animals have shown that large doses of MDMA are associated with neurodegeneration particularly affecting the terminal portions of axons and fibers; raphe cell bodies are spared (reviewed by Lyles and Cadet 2003). In rats, histological studies using the Fink–Heimer staining method have also provided evidence for the degeneration of nerve terminals in the striatum and somatosensory cortex after chronic exposure to MDMA (Commins et al. 1987). Ricaurte et al. (2000) and Callahan et al. (2001) have also provided evidence that MDMA can cause reduction of the anterograde transport of [<sup>3</sup>H] proline in ascending axons originating in the dorsal raphe nuclei, a subset of which comprise ascending 5-HT axons that project to various forebrain regions. A recent study found microtubular injury in frontal cortex after

MDMA treatment. The most apparent alteration in the ultrastructure of the labeled cortical axons of MDMA treated animals was the widespread disorganization of the microtubular system, suggesting a collapse of axonal microtubular system (with intact mitochondria) that leads to axonal swelling (Adori et al. 2011). These authors also found no signs of damage on dorsal raphe cell bodies. In nonhuman primates, reorganization of 5-HT projections has been reported to occur in the brains of nonhuman primates treated with MDMA (Ricaurte et al. 2000).

Formation of toxic MDMA metabolites which generate free radicals and associated oxidative stress and membrane damage has been proposed as causal agents for the long-term MDMA-induced neurodegeneration (reviewed in Lyles and Cadet 2003). Metabolism of MDMA results in the formation of methylenedioxyamphetamine (MDA) by *N*-demethylation and 3,4-dihydroxymethamphetamine (HHMA), the major metabolite, by *O*-demethylation. *O*-demethylation of MDA subsequently results in 3,4-dihydroxyamphetamine (HHA). HHMA and HHA are metabolized by catechol-*O*-methyltransferase (COMT) to HMMA and to 4-hydroxy-3-methoxy-amphetamine (HMA) (reviewed by Green et al. 2003). The metabolites of MDMA might induce reactive species through redox cycling. Also, DA-induced oxidative stress in 5-HT terminals, and 5-HT<sub>2A</sub> and dopamine D<sub>1</sub> receptor-mediated hyperthermia (Shioda et al. 2008) are all factors associated with MDMA neurotoxicity.

## 6 Cognition in Human MDMA Subjects

The accumulated evidence suggests that heavy MDMA use is associated with cognitive impairments and mood disturbances, in some cases lasting for months after cessation of drug intake (Baumann and Rothman 2009). Several studies have reported long-term deficits in memory and higher cognition, and other psychobiological functions in these substance abusers (Reneman et al. 2006). It is important to note that not all cognitive skills are affected and that there are also MDMA users who show normal neuropsychological functions (Parrott 2006). The role of other psychiatric disorders needs to be considered when evaluating cognitive functions in MDMA abusers. MDMA can also disrupt neuroendocrine regulation that is influenced by 5-HT levels. For example, MDMA users showed significantly reduced prolactin and cortisol responses to the serotonergic agonist d-fenfluramine in comparison with control subjects (Gerra et al. 2000).

## 7 Protective Mechanisms Against MDMA-Induced Toxicity

Preconditioning with repeated MDMA dosing appears to exert some neuroprotective effects against larger MDMA doses. For example, Bhide et al. (2009) reported that repeated exposure of adult rats to MDMA provides neuroprotection against a challenge with large doses of MDMA that were shown to deplete 5-HT and decrease in SERT immunoreactivity. Alterations in MDMA pharmacokinetics or MDMA-induced hyperthermia do not appear to contribute to the neuroprotection provided by preconditioning.

There are other agents that also provide protective effects against MDMA toxicity. Chipana et al. (2008) reported that memantine, an alpha-7 nAChR antagonist used for treatment of Alzheimer's disease, was able to provide complete protection against MDMA-induced reduction in cortical [<sup>3</sup>H]paroxetine binding sites. Fluoxetine, a SERT blocker, also provided long-lasting protection against MDMA-induced loss of SERT detected in vivo using micro-PET (Li et al. 2010).

## 8 Conclusion

In conclusion, MDMA users have been reported to suffer from cognitive and biochemical abnormalities. There is a need to correlate the changes of the serotonergic markers and other structural neuroimaging data to neuropsychological performance as well as responses to therapeutic interventions. These studies would help to provide better rationale for the development of pharmacological tools to treat METH- and MDMA-substance abusers.

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# **Glutamate as a Neurotoxin**

# Gabrielle N. Turski and Chrysanthy Ikonomidou

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## Abstract

Glutamate (Glu) is the major excitatory neurotransmitter in the mammalian central nervous system (CNS). Excessive glutamate in the extracellular space can trigger passive and active forms of neuronal death in the CNS via excessive activation of glutamate receptors. This phenomenon has been named excitotoxicity.

G.N. Turski

<sup>2</sup>nd Faculty of Medicine, Charles University, Prague, Czech Republic e-mail: ikonomidou@neurology.wisc.edu

C. Ikonomidou (⊠) Department of Neurology, University of Wisconsin, Madison, WI, USA e-mail: ikonomidou@neurology.wisc.edu

Excitotoxicity has been implicated in the pathogenesis of acute and chronic neurodegenerative disorders. However, despite overwhelming preclinical data, human clinical trials in stroke or traumatic brain injury with antiexcitotoxic compounds acting at different levels of the excitotoxic cascade have failed to provide the expected neuroprotective effect. Here we present a focused overview of excitotoxic research including physiology of glutamate receptors and intracellular pathways leading to cell death. We review evidence for involvement of excitotoxicity in human neurodegenerative disorders and the results of several clinical trials. We present the most recent trends in the development of antiexcitotoxic therapies and also briefly refer to the hypothesis that interference with the trophic actions of glutamate in the context of CNS injuries may be one of the reasons why clinical trials with antiexcitotoxic compounds have failed.

#### Keywords

Death • Excitotoxicity • Ischemia • Neuron • Neurodegenerative disorder • Stroke

# 1 Introduction

The first hint towards the neurotransmitter role of the amino acid glutamate came in 1935 from work by Krebs, who discovered that glutamate increases metabolism in the isolated retina and that it is concentrated in the cerebral gray matter (Krebs 1935). Hayashi (1952, 1958) discovered the excitatory properties of glutamate on neuronal tissue. Local administration of glutamate on the motor cortex of dogs and primates resulted in motor seizures. Curtis et al. (1959) subsequently demonstrated that glutamate and aspartate, when applied iontophoretically to the cat's spinal cord, depolarized neurons.

Today it is common knowledge that glutamate is the major excitatory neurotransmitter in the nervous system and that it is essential for learning and memory, synaptic plasticity, and neuronal survival and, in early development, for proliferation, migration, and differentiation of neuronal progenitors and immature neurons (Fonnum 1984; Guerrini et al. 1995; Ikonomidou et al. 1999; Komuro and Rakic 1993).

Compartmentalization allows that separate pools within the central nervous system supply glutamate for the metabolic machinery of the cells (metabolic pool) and for intercellular signaling (neurotransmitter pool) (Fonnum 1984). The neurotransmitter pool is located in nerve endings and releases glutamate by a calcium-dependent mechanism. Glutamate is subsequently removed by uptake into the surrounding glial cells and aminated to glutamine.

When released into the synaptic cleft, glutamate acts at the postsynaptic site on receptors (Nakanishi 1992; Hollmann and Heinemann 1994). These include the *N*-methyl-D-aspartate (NMDA), kainate, the 2-amino-3-hydroxy-5-methylisoxazole-propionate (AMPA), and the metabotropic receptors. The first three are coupled to ion

channels and are therefore called ionotropic glutamate receptors. Metabotropic glutamate receptors mediate their actions through G proteins (Hollmann and Heinemann 1994).

Excitotoxicity, a phenomenon discovered in the 1950s and extensively investigated and characterized in the 1970s and 1980s, describes a process in which excess of glutamate and other excitatory amino acid neurotransmitters causes overstimulation of neuronal glutamate receptors leading to degeneration of neurons. Lucas and Newhouse (1957) first described that systemic administration of monosodium glutamate causes degeneration of retinal ganglion cells. Olney and colleagues (Olney 1969, 1971; Olney and Ho 1970; Olney et al. 1972) subsequently observed that excitatory amino acids, when given to infant rodents and primates, cause neurodegeneration in brain areas that lack a blood–brain barrier. The ability of acidic amino acids to destroy neurons correlates with their excitatory potencies (Olney 1969, 1971; Olney and Ho 1970; Olney et al. 1972). This interaction was termed neuroexcitotoxicity. Both ionotropic and metabotropic glutamate receptors can mediate excitotoxicity (McDonald and Schoepp 1992; Schoepp et al. 1995).

# 2 Glutamate Receptors

#### 2.1 NMDA Receptors

The NMDA receptor (NMDAR) has been named for its affinity for N-methylaspartate. When activated, the NMDAR allows the influx of cations, most notably calcium. Excessive intracellular calcium concentrations activate intracellular pathways leading to physiological and pathological processes. The NMDAR exhibits a complex gating mechanism, requiring cellular depolarization in addition to the binding of ligands.

NMDA receptors represent tetrameric heteromeric subunit assemblies whose physiological and pharmacological properties depend upon their subunit composition. Three major subunit families, NR1, NR2 (A–D), and NR3 (A and B), have been cloned (Burnashev et al. 1992; Das et al. 1998; Andersen et al. 2001; Chatterton et al. 2002; Matsuda et al. 2002; Monyer et al. 1992). Most NMDA receptors in the CNS are formed from NR1 and NR2 subunits. Alternative splicing results in eight isoforms of NR1. The NR2 subunit family consists of four individual subunits termed NR2A–D (Monyer et al. 1991, 1994; Hollmann and Heinemann 1994). Different NR2 subunits result in different Ca<sup>2+</sup> permeability of the NMDA receptor channel, different gating properties, and magnesium sensitivity (Monyer et al. 1991, 1994; Ozawa et al. 1998).

Two NR1 subunits can form an ion channel which exhibits all the classical NMDAR properties, i.e., glutamate activation, magnesium block, zinc inactivation, glycine activation, interactions with polyamines, and pH sensitivity. Mayer et al. (1984) showed that depolarization of the NMDAR-expressing neuron is necessary to electrostatically remove a magnesium ion blocking the ion channel pore.

Under physiological conditions, this is accomplished by the activation of other ionotropic glutamate channels. Zinc can inhibit the NMDAR-mediated currents elicited by glutamate (Mayer and Vyklicky 1989). Glycine is a cofactor with glutamate, necessary for NMDAR channel opening (Johnson and Ascher 1987). Glycine shows different affinities at NMDA receptor subtypes. This affinity depends on the NR1 isoform and also the NR2 subunit composition of the receptor complex which allosterically influences the glycine recognition site located on the NR1 subunit (Monyer et al. 1992; Laurie and Seeburg 1994).

Polyamines can potentiate NMDAR currents at low concentrations, increase the frequency of channel opening and glycine affinity, but also reduce currents in a voltage-dependent manner at higher concentrations (Rock and Macdonald 1992; Benveniste and Mayer 1993). The polyamine/NMDAR interaction is dependent upon the subunit composition of the NMDAR channel (Williams 1997).

Protons inhibit NMDAR by interacting with the NR1 subunit on a single lysine residue (Traynelis and Cull-Candy 1990). Thus, alterations in pH can modulate NMDAR function.

NR2 subunits refine NMDAR function and modulate the characteristics of the NR1 ion channel pore. NR2B subunits also bind and link postsynaptic proteins (Sattler et al. 1999) which are parts of postsynaptic densities comprising scaffolding proteins. This allows for the coupling of intracellular enzymes (e.g., neuronal nitric oxide synthase) with ionic second messengers (e.g., Ca<sup>2+</sup> influx from NMDARs) (Aarts et al. 2002).

A third subunit associated with the NMDA receptor gene family, the NR3 subunit, is expressed in two isoforms: NR3A, which is expressed throughout the CNS, and NR3B, which is expressed primarily in motor neurons (Chatterton et al. 2002). NR1/NR3A and NR1/NR3B complexes are not activated by NMDA or glutamate, but rather elicit an excitatory Ca<sup>2+</sup> impermeant response via glycine.

# 2.2 AMPA/Kainate Receptors

AMPA receptors are comprised of four subunits, GluR1–GluR4, and require only glutamate application for activation. They are widely distributed throughout the mammalian central nervous system and mediate fast glutamatergic neurotransmission. The type of subunits within a tetrameric assembly determines biophysical and pharmacological profile of AMPA receptors. GluR1, GluR3, and GluR4 all display strong inwardly rectifying current–voltage and calcium permeability, whereas the GluR2 subunit removes calcium permeability (Hollmann et al. 1991; Burnashev et al. 1995; Geiger et al. 1995). Two splice variants for GluR subunits have been described, flip and flop; these differ in their expression between brain regions and developmental stages. AMPARs are thought to regulate the fast excitation required to remove the magnesium block of nearby NMDARs.

Other non-NMDA receptors are designated GluR5, GluR6, GluR7 (also known as GluK5–7), KA1, and KA2 (also known as GluK1–2). These form high-affinity kainate receptors (Herb et al. 1992, 1996; Hollmann and Heinemann 1994).

The properties of kainate channels are similar to AMPARs, but they are mostly impermeant to calcium ions. Kainate receptors may be localized in both the pre-synapse and post-synapse (Castillo et al. 1997; Vignes and Collingridge 1997), and application of kainate can either stimulate glutamate release (Schmitz et al. 2001) or inhibit presynaptic glutamate release (Frerking et al. 2001). Postsynaptically, kainate channels serve a similar purpose as AMPARs in alleviating magnesium block in NMDARs.

## 2.3 Metabotropic Glutamate Receptors

Metabotropic glutamate receptors (mGluRs) are single-peptide seven-transmembrane spanning proteins linked to intracellular G proteins. These are built from eight subunits, mGluR1–8. Recent evidence suggests that G-protein-independent signaling can also occur (Heuss et al. 1999). Within the mGluR family, the sequence identity varies. Classification into three subclasses (I–III) has been proposed (Nakanishi 1992; Pin and Duvoisin 1995). Within each class the amino acid sequence identity is 70 % and between classes 45 %. mGluR1 and mGluR5 belong to class I, mGluR2 and mGluR3 to class II, and mGluR4 and mGluR6–8 to class III. Different transduction mechanisms are activated by each class. Class I activates phospholipase C, increases phosphoinositide turnover and Ca<sup>2+</sup> release from internal stores, and leads to formation of diacylglycerol which then might activate protein kinase C (Abe et al. 1992; Nakanishi 1992). This group of mGluRs also modulates excitatory postsynaptic potentials at hippocampal synapses via tyrosine kinases in a G-protein-independent fashion (Heuss et al. 1999).

Classes II and III are negatively coupled to adenylate cyclase and reduce the intracellular amount of cAMP. Group II mGluRs include mGluR2s and mGluR3s. Group III mGluRs include mGluR4s, mGluR6s, mGluR7s, and mGluR8s. These mGluRs cause downstream inhibition of voltage-dependent calcium channels (Tanabe et al. 1993; Chavis et al. 1994). These receptors are found pre- and postsynaptically (Ozyurt et al. 1988; Bradley et al. 1996). Since presynaptic calcium is integral to neurotransmission, group II and III mGluRs modulate neurotransmission by functioning as autoreceptors and modulating calcium channel influxes. Group I mGluRs potentiate NMDAR-mediated Ca<sup>2+</sup> influx (Bruno et al. 1995a). mGluR2, 3 and mGluR4, 6, 7, 8 subunits reduce Ca<sup>2+</sup> influx via NMDARs (Bruno et al. 1995b).

# 3 The Neurotoxin Glutamate

The mechanisms that mediate excitotoxicity have been extensively studied in the past five decades. The acute component of glutamate toxicity is mediated through a massive influx of sodium and chloride ions into the cell (Rothman 1984, 1985; Choi et al. 1987). A second component of glutamate neurotoxicity is  $Ca^{2+}$  dependent and mediated through activation of  $Ca^{2+}$ -sensitive proteases (Choi 1987; Frandsen and Schousboe 1991, 1992; Tymianski et al. 1993a).

The concept of slow excitotoxicity arose from observations made by Novelli and coworkers showing that inhibitors of oxidative phosphorylation or of the Na<sup>+</sup>/K<sup>+</sup> pump allow glutamate or NMDA to become neurotoxic (Novelli et al. 1988). Subsequently, Zeevalk and Nicklas showed that cvanide triggers excitotoxic lesions sensitive to NMDA antagonists in the chick embryo retina (Zeevalk and Nicklas 1991). Toxicity induced by metabolic inhibition can be mimicked by membrane depolarization with potassium or by relieving the Mg<sup>2+</sup> block of the NMDA receptor (Zeevalk and Nicklas 1991). These observations were extended by findings in in vivo models showing that mitochondrial toxins (aminooxyacetic acid, an inhibitor of the maleate-aspartate shunt; 1-methyl-pyridine, a mitochondrial complex I toxin; 3-nitropropionic acid and malonic acid, inhibitors of mitochondrial complex II: the nicotinamide antagonist 3-acetylpyridine) produce axon-sparing. excitotoxic lesions in the mammalian central nervous system which can be blocked by NMDA and AMPA antagonists, glutamate release inhibitors, and prior decortication (Beal et al. 1991, 1993a, b; Greene et al. 1993; Henshaw et al. 1994; McMaster et al. 1991; Urbanska et al. 1991; Srivastava et al. 1993; Schulz et al. 1994; Wüllner et al. 1994). These observations led to the hypothesis that energy impairment at the postsynaptic site of glutamatergic synapses causes membrane depolarization, alleviation of the Mg<sup>2+</sup> block of the NMDA receptors, and heightened neuronal vulnerability towards physiological concentrations of glutamate (Beal 1992a, b; Ikonomidou and Turski 1996).

# 4 Mechanisms of Excitotoxicity

In neuronal cultures and in vivo, both active (apoptosis-like) and passive (necrosis-like) forms of cell death are seen depending on the intensity of the excitotoxic insult (Bonfoco et al. 1995; Portera-Cailliau et al. 1997). Heterogenous forms of cell death are also observed in animal models of stroke (Snider et al. 1999), hypoxia/ischemia (Fig. 1), and traumatic brain injury (Singleton and Povlishock 2004; Fig. 1). At one end of the spectrum, neurons displaying necrotic morphology are seen following intense glutamatergic insult (Bonfoco et al. 1995). The mechanisms underlying neuronal necrosis include loss of cellular homeostasis with acute mitochondrial dysfunction leading to massive energy failure. Various cell death pathways which include cysteine proteases, mitochondrial endonucleases, peroxynitrite, PARP-1, and GAPDH eventually accomplish cell death following milder glutamatergic insults. These cell death pathways operate in parallel or in sequence to orchestrate neuronal degeneration, and none of them appears to be dominant.

## 4.1 Calcium

In neuronal cultures, glutamate neurotoxicity can be separated into two components distinguishable on the basis of differences in time course and ionic dependence. The first component, marked by neuronal swelling, occurs early, is dependent on

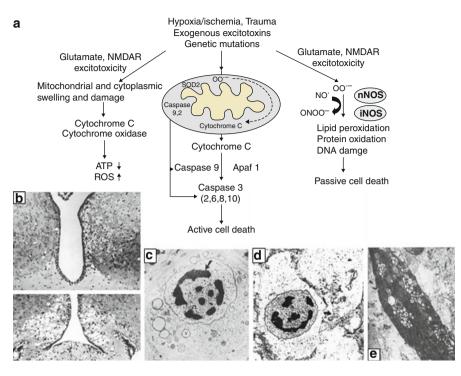


Fig. 1 Pathophysiology of cell death in the context of neurodegenerative disorders. (a) Injury generates excessive release of glutamate in the extracellular space which leads to activation of NMDA receptors and massive influx of ions into the cells. Injury also leads to generation of reactive oxygen species (ROS) within the mitochondria, which then signal the release of cytochrome c. Cytochrome c binds to Apaf-1 followed by caspase-9 to form a complex that activates caspase-3 and other caspases. The activation of the N-methyl-D-aspartate (NMDA) receptor and formation of  $O_2$ - and nitric oxide (NO) by neuronal nitric oxide (nNOS) may lead to release of cytochrome c from the mitochondria or formation of peroxynitrite (ONOO-). Resulting hydroxyl radicals can damage lipids, proteins, and DNA and lead to passive cell death. (b-e) Examples of types of neuronal degeneration seen in the brain following hypoxia/ischemia or traumatic brain injury (TBI) are shown; (b) shows neurons undergoing degeneration following hypoxia/ischemia (upper panel) in 7-day-old rats or systemic administration of monosodium glutamate in infant mice (lower panel). The pattern of degeneration in the two conditions is indistinguishable. (c) Electron micrographs depict a neuron undergoing excitotoxic death in the P7 rat brain at 4 h after hypoxia/ischemia and (d) after traumatic brain injury. Both cells display disrupted nuclear and cytoplasmic membranes, swollen mitochondria, and cytoplasmic debris. In (e) a neuron displaying dark cytoplasm filled with vacuoles is shown 4 h following traumatic brain injury in a 7-day-old rat brain. This neuron is degenerating in a more delayed fashion than shown in (b) and (c) (Adapted and modified with permission from Liebert Publications from Ikonomidou and Kaindl 2011)

extracellular Na and Cl, and is thus possibly "excitotoxic." The second component, marked by gradual neuronal disintegration, occurs late, is dependent on extracellular Ca<sup>2+</sup>, and is thus mediated by a transmembrane influx of Ca. While either component alone is ultimately capable of producing irreversible neuronal injury, the Ca-dependent mechanism predominates at lower exposures to glutamate. Glutamate exposure likely leads to a Ca influx both through glutamate-activated cation channels and through voltage-dependent Ca channels activated by membrane depolarization (Choi 1985, 1987).

Subsequent studies by Choi (1987) suggested that NMDARs may be primarily responsible for this calcium entry. Tymianski et al. (1993a) demonstrated that the path of calcium influx, and not the calcium load, was important in the NMDAR-mediated neurodegenerative process. Indeed, higher lethality can be achieved with lower calcium influxes via NMDARs compared to higher calcium influxes via other calcium-permeant channels (Sattler et al. 1998). Further work showed that NMDARs are spatially linked to neuronal nitric oxide synthase (nNOS) which can produce toxic levels of nitric oxide (NO) (Sattler et al. 1999).

The majority of intracellular calcium is sequestered into the mitochondria (Thayer and Wang 1995; Wang and Thayer 1996). Three phases of intracellular calcium concentration changes in cultured hippocampal neurons exposed to glutamate have been described: an initial phase of increased intracellular calcium lasting 5–10 min, a 2-h latent phase with normal calcium concentrations, and a gradual sustained rise in intracellular calcium associated with cell death (Randall and Thayer 1992).

Calcium activates several mechanisms to orchestrate cell death, including activation of nitric oxide synthase, proteases, and mitochondrial injury.

# 4.2 Nitric Oxide, Peroxynitrite, and Free Radicals

The findings by Dawson et al. (1991) that nitric oxide synthase (NOS) inhibitors prevent excitotoxic neuronal death in vitro provided first evidence for the role of NO in glutamate-mediated neurodegeneration. Neuronal NOS (nNOS) is the isoform causing excitotoxic injury (Dawson et al. 1996). The postsynaptic density protein of 95 kDa (PSD-95) structurally links nNOS and NMDARs (Sattler et al. 1999). The postsynaptic density protein PSD-95 binds to the C-terminus of the NR2B subunit via a PDZ1 domain and the N-terminus of nNOS via a PDZ2 domain. In addition to targeting multiple intracellular molecules (Stamler et al. 1992), NO also reacts with superoxide to form peroxynitrite, which causes cell death via protein and lipid peroxidation, protein nitration, and DNA injury (Radi et al. 1991a, b) or direct interaction with GAPDH (Hara et al. 2005). GAPDH is a ubiquitous housekeeping enzyme that participates in glycolysis. NO can nitrosylate GAPDH and bind Siah1, an ubiquitin ligase. The formed heterodimer translocates to the nucleus via Siah1 nuclear translocation domain and enhances p300/CBP-associated acetylation of nuclear proteins (Sen et al. 2009). As GAPDH is an active participant in glycolysis, it is possible that the interaction of GAPDH with nitric oxide also causes a functional loss (Molina et al. 1992) which can result in neuronal death via energy failure.

During normal cellular respiration, superoxide is produced and converted by superoxide dismutase into oxygen and hydrogen peroxide. Peroxynitrite, formed from NO and superoxide via a diffusion limited reaction (Huie and Padmaja 1993),

can cause protein nitrosylation (van der Vliet et al. 1998; Schrammel et al. 2003), lipid peroxidation (Radi et al. 1991a, b), direct DNA damage, and protein dysfunction. Peroxynitrite can also inhibit the mitochondrial electron transport chain at complex I and complex II (Radi et al. 1994), the normal function of cytochrome c (Nakagawa et al. 2001) as well as manganese and iron superoxide dismutase in scavenging superoxide (Ischiropoulos et al. 1992) via protein nitration (Yamakura et al. 1998). Peroxynitrite-mediated DNA damage can cause overactivation of the nuclear repairing enzyme poly(ADP)-ribose polymerase (PARP-1) (Zhang et al. 1994; Zingarelli et al. 1996). Evidence that free radicals play a role in glutamate excitotoxicity came from studies showing that superoxide dismutate and antioxidant compounds such as nitrone-based scavengers, free radical spin traps, and 21-aminosteroids/lazaroids confer neuroprotection (Dykens 1994; Gonzalez-Zulueta et al. 1998). Using paramagnetic resonance imaging, Lafon-Cazal et al. (1993) demonstrated a dose-dependent increase in superoxide production following increasing concentrations of NMDA. Further work in different in vitro systems demonstrated that free radical production in the context of excitotoxicity is secondary to calcium influx via NMDAR (Dykens 1994; Dugan et al. 1995; Reynolds and Hastings 1995).

# 4.3 Zinc

Zinc is present in glutamatergic neurons in a vesicle-associated pool (Martinez-Guijarro et al. 1991), which can contain up to 300  $\mu$ M concentrations of zinc (Yokoyama et al. 1986). Initial in vivo observations in kainate-treated rodents (Frederickson et al. 1989) that depletion of zinc at presynaptic terminals and simultaneous zinc accumulation in degenerating neurons occurred link zinc to glutamate neurotoxicity. Presynaptic zinc depletion, postsynaptic zinc accumulation, and neuroprotection with zinc chelation have been observed in models of cerebral ischemia (Tonder et al. 1990; Koh et al. 1996) and traumatic brain injury (Suh et al. 2000).

Zinc entry into neurons has been linked to voltage-gated calcium channels, sodium exchangers, NMDARs and AMPA/kainate receptors (Sensi et al. 1997), and TRMP7 channels (Inoue et al. 2010). Increasing intracellular zinc levels result in glycolytic dysfunction as a result of GAPDH inhibition (Sheline et al. 2000), interference with the mitochondrial electron transport chain (Nicholls and Malviya 1968), inhibition of the citric acid cycle (Brown et al. 2000), and increasing levels of reactive oxygen species (Sensi et al. 1999) via overactivation of superoxide-generating enzymes (Noh and Koh 2000) or LOX-12 (Zhang et al. 2004). Increasing levels of zinc also lead to p38 phosphorylation and ERK 1/2 activation which can trigger cell death (McLaughlin et al. 2001; Du et al. 2002).

Although the role of zinc in excitotoxic injury in animal and cell culture models is well documented, it is unclear whether modulating zinc pathways has any potential for treating human diseases.

## 4.4 Caspases and Calpains

Caspase inhibition prevents NMDA-mediated neuronal death in cerebrocortical (Tenneti et al. 1998) and cerebellar cultures (Du et al. 1997). Tenneti and Lipton (2000) described cytosolic activation of caspases at 20 min post-NMDA application with caspase activity in the nucleus at 18–24 h post-insult. Nuclear caspase activity is indicative of cleaved ICAD, a protein inhibiting the activity of caspase-3-activated DNase (CAD) and results in DNA fragmentation and cell death (Enari et al. 1998).

Brorson et al. (1995b) demonstrated modest neuroprotection in hippocampal cell cultures from NMDA insults with calpain inhibitors. The role of calpains remains less clear and may actually play a reparative role in axons with low, sublethal doses of NMDA (Faddis et al. 1997). Mu-calpain proteolytic activity is necessary in the cleavage and release of apoptosis-inducing factor (AIF) from mitochondria in a cell-free system (Polster et al. 2005). Further, calpain inhibition prevented AIF translocation and subsequent neuronal death in neuronal cultures subjected to oxygen–glucose deprivation (Cao et al. 2007), whereas other studies revealed opposite findings (Wang et al. 2009).

The existing evidence suggests that overactivation of PARP-1 secondary to an excitotoxic stimulus releases PAR polymers into the cytoplasm, which in turn activate calpains by an unclear mechanism. This, in some neuronal systems, leads to AIF release from mitochondria and subsequent cell death.

# 5 Excitotoxicity in Ischemia and Trauma

Jorgensen and Diemer (1982) first recognized a similar pattern of neuroanatomical degeneration in rats exposed to cerebral hypoxia/ischemia compared to rats treated systemically with monosodium glutamate (Olney 1969). Increased extracellular glutamate concentrations in response to ischemia were measured by microdialysis (Globus et al. 1988). In developing rats, sensitivity to NMDA neurotoxicity was found to parallel developmental vulnerability to hypoxic/ischemic injury (Ikonomidou et al. 1989), and hypoxia/ischemia was described to cause acute neuronal loss in the developing brain with ultrastructural characteristics identical to those caused by glutamate in infant rodent (Ikonomidou et al. 1989; Fig. 1). In adult global ischemia models, increases in glutamate concentrations were observed in the striatum, hippocampus, cortex, and thalamus (Globus et al. 1990). Focal ischemic models demonstrated similar patterns of elevated extracellular glutamate (Hillered et al. 1989).

Glutamate receptor antagonists were found to protect from ischemic injury. Both competitive and noncompetitive NMDA antagonists are effective in focal models of ischemia, and non-NMDA antagonists appear more effective in global cerebral ischemia models. Group I metabotropic glutamate receptor antagonists and group II agonists have been shown to be neuroprotective in gerbils subjected to transient global ischemia (Kingston et al. 1999).

Traumatic brain injury was also shown to result in marked increases in extracellular glutamate concentrations in animals (Faden et al. 1989; Nilsson et al. 1990) and in humans (Baker et al. 1993; Koura et al. 1998) and produce neuropathological changes similar to those induced by glutamate (Fig. 1). Glutamate receptor antagonists can confer neuroprotection in the context of traumatic injuries. NMDAR channel blockers reduced edema following fluid percussion brain injury in rats (McIntosh et al. 1989), whereas riluzole, a sodium channel blocker and glutamate release inhibitor, also provided histological neuroprotection (Zhang et al. 1998). Kynurenate and indole-2-carboxylic acid reduced cerebral edema and improved cognitive and motor dysfunction induced by trauma while protecting against hippocampal cell loss induced by fluid percussion injury (Hicks et al. 1994).

# 6 Excitotoxicity and Chronic Neurodegenerative Diseases

Morphological changes produced by excitotoxic compounds in the brain and spinal cord resemble in many aspects those observed in the context of chronic neurodegenerative diseases, such as Parkinson's disease, Huntington's disease, and amyotrophic lateral sclerosis. Therefore, it has been suggested that excitotoxic mechanisms may partly mediate these disorders (Beal 1992b). Especially slow excitotoxicity, triggered by the impairment of mitochondrial energy metabolism, has been implicated.

Although the genetic basis for some of these diseases has been defined, the actual mechanism(s) leading to neurodegeneration are still not understood. Slow excitotoxicity may indeed constitute a contributing factor that could be amenable to pharmacological treatment.

**Parkinson's disease (PD)** is associated with degeneration of dopaminergic neurons in the ventral mesencephalon. Clinical symptoms consist of tremor, rigidity, bradykinesia, postural deficits, and impaired gait. A gradual decline in the number of dopaminergic neurons of the substantia nigra pars compacta and dopamine content with age is common in the general population. In PD there may be an accelerated rate of cell death, so that a critical level of dopaminergic cell loss (50–70 % of neurons) is reached during normal life span and neurological symptoms of PD become obvious (Bernheimer et al. 1973; McGeer et al. 1989; Fearnley and Lees 1991). It is also possible that an exogenous insult, such as an environmental toxin, may cause a partial loss of substantia nigra dopaminergic neurons which will be followed by further "physiological" decline with age (Calne and Langston 1983). Careful neuropathological studies, serial positron emission tomography (PET), and single photon emission spectroscopy (SPECT) demonstrate a more rapid decline of dopaminergic neurons in the substantia nigra of Parkinson patients as compared to control aging subjects (McGeer et al. 1989; Fearnley and Lees 1991; Brooks 1998).

Initial observations linking excitotoxicity to PD were made by Spencer and colleagues (Spencer 1987). They reported that the excitatory amino acid  $\alpha$ -amino-beta-methylaminopropionic acid (BMAA) is linked to Guam ALS–parkinsonism dementia syndrome.

MPTP (1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine) and its metabolite MPP<sup>+</sup> (1-methyl-4-phenylpyridinium ion) have been used to model PD, since it was discovered that MPTP can cause PD in man (Davis et al. 1979; Langston et al. 1983). The first report suggesting that excitotoxicity may play a role in the pathogenesis of MPTP-induced Parkinson's disease was made by Turski and colleagues (1991), who demonstrated that neurotoxicity of MPP<sup>+</sup>, the active metabolite of MPTP, can be blocked in the substantia nigra with both competitive and noncompetitive NMDA receptor antagonists. Such results could be reproduced by other groups (Santiago et al. 1992; Brouillet and Beal 1993; Chan et al. 1993; Srivastava et al. 1993; Jones-Humble et al. 1994). Subsequent studies in primates confirmed that NMDA antagonists attenuated MPTP-induced depletion of substantia nigra dopaminergic neurons (Zuddas et al. 1992; Lange et al. 1993). Glutamate receptor blockers and neuronal NOS inhibitors have been reported to attenuate degeneration of dopamine neurons in MPTP-treated primates (Blum et al. 2001).

Considerable evidence implicates mitochondrial dysfunction in PD. Studies of the electron transport enzymes have been carried out in lymphocytes, platelets, muscle tissue, and postmortem brain tissue from PD patients (Schapira et al. 1990a, b; Di Monte et al. 1991; Hattori et al. 1991; Mann et al. 1992; Nakagawa-Hattori et al. 1992; DiDonato et al. 1993; Janetzky et al. 1994). Decrease in complex I activity was found in the brain, muscle tissue, and platelets of PD patients compared to control subjects. A nuclear magnetic resonance spectroscopy study to measure occipital lobe lactate concentration revealed a highly significant increase of lactate concentrations in PD patients with the largest increases in PD patients with dementia (Bowen et al. 1994). Similar findings indicating increased lactate levels were reported from the striatum of PD patients (Chen et al. 1994; Taylor et al. 1994).

Impaired mitochondrial function may increase vulnerability of affected neurons towards physiological concentrations of glutamate (slow excitotoxicity). Given the fact that the substantia nigra receives rich glutamatergic inputs from the neocortex and the subthalamic nucleus, contribution of slow excitotoxicity to the pathogenesis of this disease in the context of energy failure becomes a very attractive hypothesis.

Antiparkinsonian activity was observed with the noncompetitive NMDA antagonist dextromethorphan (Saez and Tanner, 1993) and in initial trials with the glutamate release inhibitor lamotrigine (Zipp et al. 1993), but these findings could not be confirmed in subsequent studies (Zipp et al. 1995; Montastruc et al. 1997). So far, the only compounds with antiglutamatergic properties that are used for the treatment of patients with PD are amantadine, memantine, and budipine (Schwab et al. 1969; Rabey et al. 1992; Adler et al. 1997). They are used to ameliorate clinical symptoms, but there is no clinical evidence indicating that they influence the course of the disease.

There have been reports that the AMPA receptor antagonist NBQX (2,3dihydroxy-6-nitro-7-sulfonyl-benzo[f]quinoxaline) improves symptoms and enhances the effect of L-3,4-dihydroxyphenylalanine (L-DOPA) in MPTP-treated monkeys (Klockgether et al. 1991; Wachtel et al. 1992), but these findings could not be replicated (Luquin et al. 1993; Papa et al. 1993; Zadow et al. 1994; Gossel et al. 1995). Due to such inconsistencies, AMPA receptor antagonists have not made their way into the clinical treatment of PD.

**Huntington's disease (HD)** is an autosomal dominant disease that gives rise to progressive, selective (localized) neural cell death associated with choreic movements and dementia. The disease is associated with increases in the length of a CAG triplet repeat present in a gene called "huntingtin" located on chromosome 4p16.3. The classic signs of Huntington's disease are progressive chorea, rigidity, and dementia, frequently associated with seizures. A characteristic atrophy of the caudate nucleus is seen radiographically. There is a prodromal phase of mild psychotic and behavioral symptoms which precedes frank chorea by up to 10 years.

The disease was initially modeled by injecting quinolinic acid (an NMDA agonist) into the striatum, leading to loss of GABAergic (GABA,  $\gamma$ -aminobutyric acid) output neurons and sparing interneurons containing somatostatin-NPY (Schwarcz and Kohler 1983; Schwarcz et al. 1983). This neuropathological pattern resembles that seen in Huntington's disease brains (Schwarcz et al. 1983; Beal et al. 1986). Based on these findings, an excitotoxicity hypothesis for the pathogenesis of Huntington's disease was formulated.

Administration of the mitochondrial toxin 3-nitropropionic acid (3-NP) to rodents produces pathology resembling Huntington's disease (Brouillet et al. 1995; Palfi et al. 1996; Borlongan et al. 1997). Interestingly, striatal toxicity of 3-NP can be ameliorated by NMDA receptor antagonists, such as memantine (Wenk et al. 1996), and glutamate release inhibitors, such as riluzole (Guyot et al. 1997). Similarly, striatal toxicity of malonate, a complex II–III inhibitor, was prevented by NMDA receptor antagonists MK-801 and memantine and by the glutamate release inhibitor lamotrigine (Greene and Greenamyre 1995). Thus, it has been postulated that mitochondrial dysfunction in Huntington's patients may be a trigger that constitutes striatal neurons prone to excitotoxicity.

The mechanism through which the widely expressed mutant HD gene mediates slowly progressing striatal neurotoxicity is unknown. Glutamate receptor-mediated excitotoxicity has been hypothesized to contribute to HD pathogenesis. However, inconsistent with this hypothesis, Hansson et al. (1999) showed that transgenic HD mice expressing exon 1 of the human HD gene with an expanded number of CAG repeats are strongly protected from acute striatal excitotoxic lesions. Intrastriatal infusions of quinolinic acid, the agonist of the NMDAR, caused massive striatal neuronal death in wild-type mice, but no damage in transgenic HD littermates. The remarkable neuroprotection in transgenic HD mice occurred at the stage when they had not developed any neurologic symptoms caused by the mutant HD gene. At this stage, there was no change in the number of striatal neurons and astrocytes in untreated transgenic mice, although the striatal volume was decreased by 17 %. Hansson et al. (1999) proposed that the presence of exon 1 of the mutant HD gene induces profound changes in striatal neurons that render these cells resistant to excessive NMDA receptor activation.

The antiglutamatergic compound remacemide, tested in a double-blind placebocontrolled study in HD patients, showed only a trend towards symptomatological improvement (Kieburtz et al. 1996). Ketamine failed to show any symptomatological improvement (Murman et al. 1997). Thus, there is so far no evidence that symptomatological improvements should be expected by the use of antiglutamatergic compounds in HD patients.

Alzheimer's disease (AD) is by far the most common cause of dementia. Terry and Davies (1980) recommended the term senile dementia of the Alzheimer type (SDAT). Senile (neuritic) plaques and neurofibrillary tangles comprise the major neuropathological lesions in AD brains. Neuritic plaques are spherical lesions and are found in moderate or large numbers in limbic structures and association neocortex. They contain extracellular deposits of amyloid- $\beta$  protein. Degenerating axons and dendrites are present within and around the amyloid deposits (Terry and Davies 1980; Selkoe 1999).

There are some indications that glutamate and the mechanism of excitotoxicity may be involved in pathogenesis and progression of AD. Constituents of senile plaques stimulate production by microglia of an unknown NMDA agonist (Giulian et al. 1995) as well as production of nitric oxide (NO), known to enhance glutamate release and inhibit its uptake (Lees 1993; Goodwin et al. 1995).  $\beta$ -Amyloid enhances glutamate toxicity (Brorson et al. 1995a; Mattson and Goodman 1995), enhances depolarization dependent glutamate release (Arias et al. 1995), and inhibits its glial uptake (Harris et al. 1996).

There is a decrease in astroglial glutamate EAA2 carrier in the frontal cortex of AD patients. Colocalization of glutamate neurons and neurofibrillary tangles was reported in AD brains (Braak et al. 1993; Francis et al. 1992). Finally, intracerebroventricular administration of  $\beta$ -amyloid produced long-lasting depression of EPSP that was prevented by the NMDA antagonist CPP (Cullen et al. 1996), whereas subcutaneous administration of memantine prevented pathological alterations in the hippocampus produced by direct injection of  $\beta$ -amyloid.

**Amyotrophic lateral sclerosis (ALS)** is the most common motor neuron disease in adults. It is characterized by selective degeneration of upper and lower motor neurons, progressive weakness and paralysis of all muscles, and spasticity (Cleveland and Rothstein 2001). Over 90 % of cases occur sporadically; in 5–10 % of cases the disease is inherited in a dominant mode (familial ALS).

Several pathogenetic mechanisms have been suggested. These include oxidative damage, neurofilament disorganization and subsequent axonal damage, toxicity from intracellular aggregates, and excitotoxicity (Cleveland and Rothstein 2001).

There is unquestionable and landmark evidence that familial ALS and some forms of sporadic ALS are caused by mutations in the Cu/Zn superoxide dismutase (SOD1) gene (Gaudette et al. 2000; Andersen 2001; Andersen et al. 2001). Multiple mutations have been identified so far and almost all provoke a dominantly inherited disease.

Debate on the involvement of excitotoxicity in ALS is based on studies suggesting that the metabolism of glutamate is abnormal in patients with ALS. Rothstein et al. (1992) postulated that the high-affinity glutamate transporter is the site of the defect at least in some forms of the disease. The inhibition of glutamate transport is toxic to neurons, due to persistent elevation of extracellular

glutamate. A possible mechanism for the elevated cerebrospinal fluid concentrations of glutamate and aspartate in patients with ALS could be deficient transport into cells. Studying synaptosomes from neural tissue obtained from 13 patients with ALS as well as from controls, Rothstein et al. (1992) found that ALS patients showed a marked diminution in the maximal velocity of transport for high-affinity glutamate uptake in synaptosomes from the spinal cord, motor cortex, and somatosensory cortex, but not in those from visual cortex, striatum, or hippocampus. This diminution was the result of a pronounced loss of the astroglial EAAT2 protein (Rothstein et al. 1995). Transport of other molecules (gamma-aminobutyric acid and phenylalanine) was normal in patients with ALS.

The fact is that in about one third of ALS patients, glutamate concentrations are increased in the cerebrospinal fluid (CSF) (Shaw et al. 1995), whereas intracellular glutamate concentrations are decreased (Perry et al. 1987; Young 1990; Ludolph et al. 1988). CSF from ALS patients is toxic to hippocampal neurons via stimulation of AMPA receptors (Couratier et al. 1993). The glutamate release inhibitor riluzole attenuated damage induced by ALS CSF (Couratier et al. 1994; Terro et al. 1996). These findings, along with experimental evidence implicating selective vulnerability of spinal motoneurons to AMPA/kainate-receptor agonists (Hugon et al. 1989; Ikonomidou et al. 1996; Williams et al. 1997), implicate that glutamate acting on AMPA/kainate receptors may be pathogenetically involved in motor neuron degeneration in ALS.

Excitotoxicity has been implicated in two other neurological disorders that resemble ALS, the Guam ALS–parkinsonism dementia syndrome and a disease described in the Kii peninsula of Japan. These two disorders have been linked to food excitotoxins (Spencer et al. 1986, 1993).

Along with the finding that SOD1 mutants, when expressed in mice, lead to functional loss of EAAT2 (Nagano et al. 1996), it seems likely that deficient glutamate uptake may contribute to the pathogenesis of some forms of ALS. Riluzole, a glutamate release inhibitor, achieved a modest effect at increasing survival in two independent trials (Bensimon et al. 1994; Lacomblez et al. 1996) and has been approved for the treatment of ALS, although patients do not experience slowing in disease progression.

# 7 Clinical Experience with Drugs Targeting Excitotoxicity in Stroke and Traumatic Brain Injury

Overall clinical experience has been extremely disappointing. Even though animal models demonstrate dramatic histological and behavioral neuroprotection, no pharmacological intervention targeting excitotoxicity has yet been shown to provide benefit in humans (Muir 2006).

Noncompetitive NMDAR antagonists have been shown in animal models to provide histological and behavioral neuroprotection following focal ischemia (Block and Schwarz 1996; Ozyurt et al. 1988; Seif et al. 1990; Steinberg et al. 1988). Cerestat (CNS 1102) was tested in a multicenter trial in 628 stroke patients

with low-dose, high-dose, and placebo treatments. The trial was stopped because of lack of effect and a trend towards higher mortality with high-dose Cerestat compared to placebo (Albers et al. 2001). Memantine has shown a much better side effect profile in humans (Areosa et al. 2005) and is currently approved for use only in moderate to severe Alzheimer's disease (Reisberg et al. 2003). No stroke trials with memantine have been performed.

The competitive NMDA antagonist Selfotel (CGS 19755) was shown to reduce hippocampal damage in gerbil models of ischemia (Boast et al. 1988). Phase III trials in stroke and in severe head injury were abandoned due to concerns about higher mortality in the treatment arms (Davis et al. 2000; Morris et al. 1999).

The glycine antagonist Gavestinel reached phase III clinical trials, but no difference in morbidity or mortality rates following treatment in stroke was evident (Lees et al. 2000). Ifenprodil and eliprodil, drugs binding at the NMDAR polyamine site, underwent phase III clinical trials but showed no efficacy.

AMPA receptor antagonists initially appeared more effective and showed a longer time-window in preventing neuronal loss in animal models of global ischemia (Sheardown et al. 1990; Buchan et al. 1991). YM872 demonstrated substantial neuroprotective effects in animal models of focal ischemia (Takahashi et al. 1998) and traumatic brain injury models (Furukawa et al. 2003), but phase III trials in ischemic stroke showed no efficacy in 2006.

Lamotrigine inhibits presynaptic voltage-dependent sodium channels and reduces overall excitability and neurotransmitter release (Leach et al. 1986). In animal models, lamotrigine failed to provide neuroprotection in focal ischemia (Tymianski et al. 1993a, b) but showed a reduction of CA1 hippocampal loss of 50 % following global ischemia (Crumrine et al. 1997). BW619C89, a derivative of lamotrigine, underwent phase II trials in 16 of 21 stroke patients (Muir et al. 2000). Clinical development was halted in 2001. Riluzole has also been shown to reduce infarct volumes and improve behavioral outcomes in ischemic (Malgouris et al. 1989; Wahl et al. 1993) and traumatic brain injury (Wahl et al. 1997; Zhang et al. 1998), but no clinical trials for stroke or traumatic brain injury have been performed. Riluzole is licensed for use in ALS (Miller et al. 2007).

# 8 Clinical Trials with Free Radical Scavengers

The nitrone-based antioxidant NXY-059 was tested in stroke. While the SAINT1 trial examining the efficacy of NXY-059 in clinical stroke was promising (Lees et al. 2000), the SAINT2 trial showed no clinical benefit with treatment (Shuaib et al. 2007). Human clinical trials with the 21-aminosteroid Tirilazad<sup>®</sup> in head injury resulted in decreased mortality only in males with severe head injury patients and subarachnoid hemorrhages (Marshall et al. 1998). Subsequent trials with Tirilazad in the treatment of subarachnoid hemorrhages (Lanzino and Kassell 1999; Lanzino et al. 1999) showed a reduction in mortality only with the highest grades on arrival, but meta-analysis of Tirilazad in the treatment of subarachnoid aneurysmal hemorrhage demonstrated no clinical outcome differences, but reduced

symptomatic vasospasm (Jang et al. 2009). A Cochrane Review demonstrated increased death or disability in stroke victims treated with Tirilazad compared to placebo in a review of six clinical trials without any statistically significant difference in overall mortality (Bath et al. 2001). Studies in ischemic stroke using Tirilazad have since been abandoned.

In a large multicenter stroke trial, the free radical trapping agent NXY-059, when given within 6 h after the onset of acute ischemic stroke, significantly improved the primary outcome (reduced disability at 90 days), but it did not significantly improve other outcome measures, including neurologic functioning as measured by the NIHSS score (Lees et al. 2006).

Ebselen is a glutathione peroxidase which can interact with peroxynitrite and inhibit enzymes involved in inflammation (Schewe 1995). Clinical trials in Japan demonstrated no significant differences in a 3-month clinical outcome in patients with complete middle cerebral artery occlusion (Yamaguchi et al. 1998) and no significant improvement in clinical outcome despite a reduction in infarct size (Ogawa et al. 1999).

A novel class of drugs named membrane-associated guanylate kinase (MAGUK) inhibitors are currently undergoing phase II clinical trials. The MAGUK inhibitor NA1 uncouples nNOS and NMDARs and has demonstrated neuroprotection in various animal models of stroke (Aarts et al. 2002; Sun et al. 2008).

# 9 Conclusion

Despite the overwhelming evidence that a variety of neurotoxic mechanisms triggered by excessive glutamate concentrations mediate brain injury in human neurological disorders, all treatment attempts with antiglutamatergic agents in humans have failed. One hypothesis for this failure is that glutamate has a dual function in the injured brain, a toxic one and a trophic one, and that the latter has been ignored.

Synaptic transmission mediated by NMDA receptors is essential for neuronal survival. Blockade of NMDA receptors for only a few hours triggers apoptosis in the developing brain (Ikonomidou et al. 1999; Pohl et al. 1999). NMDA antagonists, when administered during a critical period following traumatic brain injury or in the context of slowly ongoing neurodegeneration, markedly exacerbate damage in adult brain (Ikonomidou et al. 2000).

The Ca<sup>2+</sup> pool in the immediate vicinity of synaptic NMDA receptors is the on switch for extracellular signal-regulated kinase (ERK1/2)-mediated synapse-tonucleus signaling (Hardingham et al. 2001). One important function of this Ca<sup>2+</sup> microdomain near NMDA receptors is to prolong phosphorylation of the transcription factor cyclic AMP response element-binding protein (CREB) induced by synaptic stimuli, thereby enhancing CREB-mediated gene expression. CREB controls transcription of prosurvival genes such as brain-derived neurotrophic factor (BDNF), vasoactive intestinal peptide (VIP), bcl-2, and mcl-1 (Fink et al. 1991; Tao et al. 1998; Riccio et al. 1999). It is therefore postulated that survival-promoting properties of NMDAR activation derive from the transcription of such prosurvival genes. Neurons surviving ischemic insults (penumbra) demonstrate elevated concentrations of BDNF, bcl-2, and activated CREB, suggesting sustained induction of prosurvival signals (Chen et al. 1995; Tanaka et al. 1999, 2000; Walton and Dragunow 2000). From these findings the logical conclusion is that suppression of NMDA receptor-initiated survival-promoting signals with NMDA antagonists may facilitate death of such cells. Blockade of NMDA-mediated synaptic transmission must therefore be detrimental in situations, in which support by endogenous measures is required, as it happens after stroke or traumatic brain injury or in chronic neurodegenerative disorders (Ikonomidou and Turski 2002).

The long-term elevation of glutamate concentration in humans (days to weeks) after traumatic brain injury has been considered neurotoxic and interpreted as an opportunity for delayed therapy with NMDA antagonists (long time-window for therapy initiation). However, this interpretation may be wrong and such mild elevations of glutamate concentration may represent a self-defense mechanism of the injured brain.

Glutamate may indeed be involved in the initial acute neurodestructive phase after traumatic or ischemic injury, but after this period, it assumes its normal physiological functions, which include promotion of neuronal survival. The hypothesis was that mild increases of glutamate concentrations starting shortly after brain injury are necessary for supporting survival of neurons endangered to die after the insult and for maintaining their physiological functions. Conforming with this hypothesis, it was possible to demonstrate that NMDA antagonists are detrimental to neurons subjected to traumatic brain injury when administration starts after the initial profound elevation of extracellular glutamate concentrations has subsided, that is, 1-7 h after trauma (Ikonomidou et al. 2000). Antecedent treatment with NMDA antagonists in this same setting prevented traumatic neuronal death. These findings strongly argue for the hypothesis that glutamate indeed kills neurons over a brief period after injuries but starts to facilitate repair immediately thereafter. In contrast to its killer function, repair mediated by glutamate appears to be a long-lasting phenomenon. This is in obvious agreement with physiological function of glutamate in the nervous system during development.

Thus, to be successful in preventing excitotoxic brain injury, we would need to specifically target the neurodestructive pathways activated by glutamate while preserving the ones that promote survival. One promising strategy is in the horizon. PSD-95 inhibitors have been shown to confer neuroprotection in stroke models in rodents (Aarts et al. 2002; Sun et al. 2008). The 20-mer peptide Tat-NR2B9c, which acts by perturbing the protein–protein interactions of PSD-95, a synaptic protein that links NMDA receptors to neurotoxic signaling pathways, was most recently shown to confer histological and behavioral neuroprotection in gyrencephalic primates (Cook et al. 2012). This work rejuvenates hopes that more sophisticated and exclusive blockade of neurotoxic but not trophic mechanisms triggered by glutamate may one day provide a mode of treatment that will enable us to limit brain injuries in human patients.

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# Domoic Acid as a Neurotoxin

# Anabel Pérez-Gómez and R. Andrew Tasker

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#### Abstract

Domoic acid (DOM) is a naturally occurring excitatory amino acid with structural similarity, and similar but not identical pharmacological profile, to kainic acid. DOM is most commonly associated with toxic blooms of marine phytoplankton resulting in contamination of shellfish as well as other species. This brief review summarizes the known toxicological properties of DOM both in vitro and in vivo in a variety of model systems. This chapter also summarizes information on clinical cases of intoxication in both wildlife and humans, as well as highlighting the growing evidence that DOM is a potent neurodevelopmental toxin with relevance to both food safety issues and the etiology of neurological diseases.

A. Pérez-Gómez • R.A. Tasker (⊠)

Department of Biomedical Sciences, University of Prince Edward Island, Charlottetown, PEI, Canada

e-mail: aperez@upei.ca; tasker@upei.ca

#### Keywords

Amnesic shellfish toxin • Excitatory amino acid • Excitotoxicity • Glutamate • Marine toxin

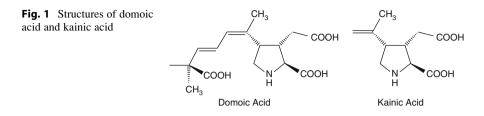
# 1 Chemistry, Physicochemical Properties, and Pharmacology

Domoic acid is a naturally occurring excitatory amino acid (EAA) that has been linked to numerous outbreaks of toxicity in humans (Teitelbaum et al. 1990; Cendes et al. 1995) as well as a variety of other species (for review see Doucette and Tasker 2008; Lefebvre and Robertson 2010). The parent compound is a tricarboxylic amino acid produced by certain marine organisms, the best characterized of which are the red alga *Chondria armata* and the planktonic diatoms of *Nitzschia spp*. (for review see Bates and Horner 1998; Bates and Trainer 2006).

Chemically domoic acid is (2S,3S,4S)-2-carboxy-4-1-methyl-5(R)-carboxyl-1 (Z)-3(E)-hexadienyl pyrrolidine-3-acetic acid (C<sub>15</sub>H<sub>21</sub>NO<sub>6</sub>) and has an anhydrous molecular weight of 311.14, a melting point of 215–216 °C (dihydrate), and a UV (ethanol) absorption spectrum with a maximum of 242 nm although this is pH dependent. Structurally domoic acid is very similar to another known toxin, kainic acid, and both are analogues of the amino acids glutamate and proline. The structures of domoic and kainic acid are shown in Fig. 1.

In addition to the parent compound, a number of isomers of domoic acid have been identified. Isodomoic acids A, B, and C are present in small amounts in the red alga *Chondria armata* (Maeda et al. 1986) as are isodomoic acids G and H (Zaman et al. 1997). The isomers isodomoic acids D, E, and F as well as the 5' epimer have been identified in small amounts from both plankton cells and shellfish tissue (Wright et al. 1990; Walter et al. 1994) although it seems likely that these isomers form on exposure to UV light rather than being metabolic products produced by the plankton (Quilliam 2003). Although the binding of domoic acid isomers in vitro to kainate receptors (Rs) was reported a number of years ago (Hampson et al. 1992), there have been few investigations to date of the pharmacology and toxicology of domoate isomers, with the exception of four publications out of New Zealand (Sawant et al. 2007, 2008; Munday et al. 2008; Sawant et al. 2010).

Both DOM and its analogue kainic acid are excitotoxic and mediate their toxic effects through activation of glutamate receptors (GluRs) with the participation and co-activation of both *N*-methyl-D-aspartate (NMDA) receptors and non-NMDA-R subtypes (AMPA [alpha-amino-3-hydroxy-5-methylisoxazole-4-propionic acid]/ kainate) (Novelli et al. 1992; Larm et al. 1997; Berman and Murray 1997; Berman et al. 2002; Hampson and Manalo 1998; Clements et al. 1998; Tasker and Strain 1998; Tasker et al. 2005). While DOM stimulates AMPA/kainate-Rs (AMPA/KA-Rs) directly, NMDA-R activation appears to be a consequence of either AMPA/ KA-R-mediated stimulation of EAA efflux (Novelli et al. 1992; Berman and Murray 1997; Clements et al. 2002) and/or to reduce tonic



inhibition by gamma-aminobutyric acid (GABA) as a result of activation of presynaptic kainate-Rs (KA-Rs) on GABAergic interneurons (Duran et al. 1995; Cunha et al. 1997; Rodriguez-Moreno et al. 2000). Although similar in many respects, DOM and kainic acid are pharmacologically dissociable in some systems because DOM interacts preferentially with the so-called low-affinity kainate subunits designated GluK1 and GluK2 (previously referred to as GluR5 and GluR6) (Verdoorn et al. 1994; Tasker et al. 1996). The complexity of DOM pharmacology combined with the relative lack of knowledge about the physiological properties of KA-Rs probably explains some of the discrepancies reported in the literature, particularly between in vitro and in vivo preparations (see below).

# 2 DOM Toxicity In Vitro

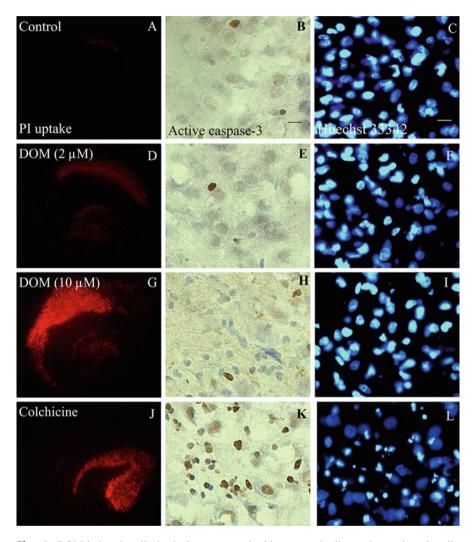
DOM-induced neurotoxicity has been evaluated in a variety of in vitro systems including immature and mature primary cultures of neurons from cerebellum, cortex, and hippocampus, organotypic hippocampal cultures, and neuroblastoma cell lines. DOM-induced neuronal death is dependent on both exposure duration and concentration, with an EC<sub>50</sub> reported to range from 3 to 10  $\mu$ M after 24 h exposure in cerebellar granule cell cultures from both rat and mouse (Novelli et al. 1992; Fernandez-Sanchez and Novelli 1993; Larm et al. 1997; Giordano et al. 2006; Perez-Gomez et al. 2010). The EC<sub>50</sub> values in cortical neurons range from 4.2 µM in rat (Qiu et al. 2006) to higher than 20 µM in mouse (Jensen et al. 1999) after 24 h treatment, and in mixed hippocampal cultures, an EC<sub>50</sub> of 7.5  $\mu$ M has been reported (Roy and Sapolsky 2003). An EC<sub>50</sub> of 3 µM was obtained for the NG108-15 neuroblastoma cell line (Canete and Diogene 2008), and finally, the treatment of organotypic hippocampal cell cultures (OHSCs) with DOM resulted in selective but not specific toxicity to CA1 pyramidal neurons with an EC<sub>50</sub> of 3.5  $\mu$ M after 24 h of exposure (Perez-Gomez and Tasker 2012) and 6  $\mu$ M after 48 h exposure (Jakobsen et al. 2002). DOM effects in vitro confirm results obtained in vivo showing that DOM is 3-20-fold more potent than KA itself (Stewart et al. 1990; Tasker et al. 1991).

The mechanism of action responsible for DOM toxicity in vitro appears to be more complex than would be predicted from the compound's pharmacological profile as an AMPA/KA-R agonist. DOM in vitro elicits a non-desensitizing response via activation of low-affinity AMPA-Rs (Diaz-Trelles et al. 2003), while activation of KA-Rs does not seem to play a crucial role (Berman and Murray 1997; Jakobsen et al. 2002; Diaz-Trelles et al. 2003; Gressens et al. 2005). DOM has also been shown to evoke the release of both glutamate and aspartate in cerebellar granule cells (CGCs) (Berman and Murray 1997) as well as in isolated chick retina explants (Nduaka et al. 1999). This appears to occur via a non-vesicular mechanism that is dependent upon reversal of the high-affinity glutamate transporter and cell swelling and, therefore, synergistically potentiates glutamate-/aspartate-mediated neurotoxicity (Berman and Murray 1997). Pharmacologically, both NMDA-R and non-NMDA-R have been implicated in the toxic actions of DOM, in that DOMinduced injury seems to be mediated in part by NMDA-Rs that are activated secondarily as a consequence of AMPA-R-mediated stimulation of EAA release (Berman and Murray 1997). Accordingly, DOM has been shown to act synergistically with NMDA in producing acute cytotoxicity in CGCs (Novelli et al. 1992; Berman and Murray 1996).

In vitro studies have shown increased DOM toxicity with increasing maturation of the central nervous system (Qiu et al. 2006; Hogberg and Bal-Price 2011). This could be due to lower functional expression of the GluRs in the younger cultures. It has also been proposed that more calcium-permeable AMPA-Rs or AMPA-R isoforms could be activated in the mature cultures in comparison to the younger cultures (Qiu et al. 2006), causing glutamate release that subsequently could activate the NMDA-R, promoting neurotoxicity (Berman and Murray 1997; Berman et al. 2002).

Application of DOM produces a rapid and concentration-dependent increase in intracellular calcium in a variety of neuronal preparations, including embryonic brain stem cells (Konig et al. 1994), dorsal root ganglion neurons (Joseph et al. 2011), cultured neurons of chick embryonic retina (Duran et al. 1995), and primary cultures of cortical (Beani et al. 2000), hippocampal (Xi and Ramsdell 1996), cerebellar granule (Perez-Gomez et al. 2010), and Purkinje neurons (Gruol et al. 1997). Vale-Gonzalez et al. (2006) described a prominent intracellular acidification resulting from a rise in intracellular calcium. Some studies have proven that DOMstimulated increases in cytosolic-free calcium are dependent upon the activation of L-type voltage-sensitive calcium channels (VSCC) following activation of non-NMDA-R (Xi and Ramsdell. 1996; Berman et al. 2002). However, the contribution of a calcium increase through VSCC to DOM toxicity remains controversial as some data suggested that calcium may reduce rather than potentiate the nondesensitizing AMPA-R response to DOM in CGCs, thereby decreasing neurotoxicity as well as the generation of reactive oxygen species (Fernandez-Sanchez and Novelli 1993; Diaz-Trelles et al. 2003; Perez-Gomez et al. 2010).

In some in vitro preparations, DOM-induced neuronal cell death appears to be primarily necrotic in nature. No signs of the classical caspase-dependent apoptotic pathway were detected in organotypic hippocampal slices using different concentrations and time of exposures to DOM (Perez-Gomez and Tasker. 2012) (Fig. 2). In mixed hippocampal cell cultures, DOM neurotoxicity was also found to be necrotic when examined both biochemically and morphologically (Roy and Sapolsky. 2003), and protection afforded by two virally derived caspase inhibitors (p35 and crmA) was ascribed to their ability to sequester free radicals and to stabilize the mitochondrial



**Fig. 2** DOM-induced cell death in organotypic hippocampal slice cultures is primarily necrotic. Propidium iodide exclusion ( $\mathbf{a}$ ,  $\mathbf{d}$ ,  $\mathbf{g}$ ) demonstrates that cell death in OHSCs is concentration dependent but is not accompanied by an increase in the apoptosis marker activated caspase-3 ( $\mathbf{e}$ ,  $\mathbf{h}$ ) relative to control ( $\mathbf{b}$ ) or a change in the intensity of Hoechst 33342 staining for condensed chromatin ( $\mathbf{f}$ ,  $\mathbf{i}$ ) relative to control ( $\mathbf{c}$ ). In contrast administration of colchicine as a positive control results in comparable cell death ( $\mathbf{j}$ ) accompanied by changes in both caspase-3 and Hoechst 33342 staining ( $\mathbf{k}$ ,  $\mathbf{l}$ )

membrane potential, respectively. However, acute in vitro studies in pure mouse CGCs have shown that DOM can cause either necrotic or apoptotic cell death, depending on its concentration. A high concentration of DOM (1 h, 10  $\mu$ M) activates both the AMPA/KA-R and NMDA-R which causes a rapid accumulation of intracellular calcium; promotes GSH efflux and a concomitant decrease in

intracellular GSH, inducing an increased lipid peroxidation; and leads to necrotic neuronal death (Giordano et al. 2006). On the other hand, AMPA/KA-Rs (not NMDA-R) have been shown to mediate low-concentration ( $<0.1 \mu$ M) DOM-induced apoptosis (Giordano et al. 2007) with DOM inducing oxidative stress, leading to mitochondrial dysfunction and activation of caspase-3. Similar results were obtained using the motor neuron-like cell line NSC34, where DOM dramatically increased reactive oxygen species production, reduced mitochondrial function and cell viability, and was accompanied by an increase of cell apoptosis (Xu et al. 2008). A moderate increase in degraded  $\alpha$ -spectrin concentration was found in organotypic rat brain slices (Erin and Billingsley 2004). The importance of oxidative stress in DOMinduced apoptosis was confirmed in mouse CGCs by the fact that activation of muscarinic receptors provides protection against DOM toxicity in part by an increase in the antioxidant capacity of the cells (Giordano et al. 2009). In addition, and by using the same experimental system (mouse CGCs), Giordano et al. (2008) have shown that DOM also causes an increase in the phosphorylation of p38 and Jun N-terminal (JNK) kinases, which are preferentially activated by cell stress-inducing signals (Raman et al. 2007).

While many reports have described the effects of high concentrations of DOM on neuronal systems in vitro, the effects of low-concentration DOM have also been tested in a variety of in vitro systems, leading to interesting results. Indeed, longterm exposure of primary cortical neurons to low-concentration DOM (50 nM, 4 weeks), which acutely did not induce any effects on electrical activity in the mature cultures, significantly increased the spontaneous activity of neurons when compared to non-treated cultures (Hogberg et al. 2011). The authors concluded that DOM should be recognized as a developmental neurotoxicant as the induced changes in electrical activity could lead to serious disturbance of overall neuronal activity in the developing brain. This conclusion is consistent with the findings of a number of in vivo studies in perinatal animals (see subsequent section) as well as with a recent report by Perez-Gomez and Tasker (2012) in which organotypic hippocampal slice cultures were exposed to concentrations below those required to induce permanent neurotoxicity. In this study, low-concentration DOM produced an increase of proliferation and neurogenesis in the dentate gyrus and CA1 subfields that may contribute to the development of neuroplastic changes and abnormal circuits relevant to disease. Along similar lines, Perez-Gomez et al. (2010) demonstrated that mild, nontoxic exposures to the Na<sup>+</sup>/K<sup>+</sup>-ATPase inhibitor palytoxin synergistically sensitized the vulnerability of CGCs to normally nontoxic concentrations of DOM, leaving NMDA receptor-mediated excitotoxic response unaltered. The results obtained demonstrate not only a crucial role for Na<sup>+</sup>/K<sup>+</sup>-ATPase activity in determining neuronal vulnerability to DOM-mediated excitotoxicity, but, as palytoxin and DOM are both seafood contaminants found in geographically overlapping regions, the results raise reasonable concern about possible risks for human health associated to the ingestion of low (i.e., below regulatory limits), subtoxic amounts of these phycotoxins in food.

A few studies have been conducted exploring the possibility that DOM affects not only neurons but also other cells of the central nervous system such as astrocytes or microglia. It has been shown that exposure of astrocytic cultures to DOM did not induce any cell death (Novelli et al. 1992; Ross et al. 2000; Giordano et al. 2006), but, to a certain extent, it changed some glial functions. Injury to the astrocytes was characterized by ultrastructural changes and altered expression of chemokines/cytokines, tyrosine kinases (Trk), and apoptotic genes (Gill et al. 2008). Changes in the mRNA expression of glial fibrillary acidic protein (GFAP) (downregulation) and S100 $\beta$  (upregulation) (Hogberg and Bal-Price 2011) as well as effects on membrane function of rat glial cells (Liu et al. 2008) and modest increases in intracellular calcium (Giordano et al. 2006) have been reported after DOM treatment. Involvement of glial cells could enhance DOM-induced neurotoxicity as both glutamate uptake and its metabolism to glutamine were significantly inhibited (Ross et al. 2000).

In accordance with the results in astrocytes, DOM does not seem to induce toxicity to microglial cells in vitro (Mayer et al. 2001). The hypothesis that DOM might lead to the activation of microglia has been tested as well, and results seem to depend on the duration of the exposure as well as on the type of culture used. In organotypic hippocampal slice cultures, low-dose DOM insult ( $2 \mu M$ , 24 h) induced a significant increase in the number of reactive microglial cells 7 days after treatment (Perez-Gomez and Tasker 2012). On the other hand, DOM treatment (4–24 h, 1 mM) does not appear to activate neonatal rat microglial cultures and the concomitant expression and release of pro- and anti-inflammatory mediators (Mayer et al. 2007).

## 3 DOM Toxicity In Vivo

#### 3.1 Absorption, Distribution, and Elimination

DOM is usually administered parenterally (i.p. or s.c.) in most experimental paradigms, but the more common route of exposure for intoxication of wildlife and humans (see Sects. 4.1 and 4.2 below) is by oral ingestion. Surprisingly, however, there have been very few objective studies on the absorption of this toxin following oral administration. Based on these limited observations, it appears that DOM absorption from the GI tract is erratic and bioavailability is approximately 5–10 % whether studied in mice and rats (Iverson et al. 1989, 1990) or in cynomolgus monkeys (Iverson et al. 1990; Tryphonas et al. 1990b; Truelove et al. 1996). Bioavailability following parenteral administration (i.p. or i.v.) on the other hand appears to be complete.

Once absorbed DOM appears to be largely restricted to the vascular compartment with an interpolated apparent steady-state volume of distribution (Vd<sub>ss</sub>) of approximately 0.20 l/kg in all species studied to date (Suzuki and Hierlihy 1993; Truelove and Iverson 1994). This is consistent with the very hydrophilic nature of the molecule, as is the comparatively poor blood–brain barrier penetration of DOM in normal adult rats, where mean transfer constants ranging from 1.60 to  $1.82 \text{ ml/g/s} \times 10^6$  have been reported (Preston and Hynie 1991) (in the same study sucrose transfer constants ranged from 1.00 to 1.24). These data argue against the existence of a carrier protein and suggest that the highly charged state of DOM at physiological pH results in poor CNS penetration kinetics, which seems at variance with the known CNS toxicity associated with DOM (see below). On closer examination, however, the data suggest that DOM is an extremely potent neurotoxin within the CNS, an assumption that is confirmed by studies reporting extremely low  $ED_{50}$  values when DOM is administered directly into the CNS via intracerebroventricular (i.c.v.) injection (Sawant et al. 2008; Vranyac-Tramoundanas et al. 2011) or is administered to very young rats prior to blood–brain barrier closure (Xi et al. 1997; Doucette et al. 2000).

DOM appears to be eliminated almost exclusively by renal excretion. Preston and Hynie (1991) reported that a single i.v. dose of radiolabelled DOM was almost completely eliminated from the serum of intact rats within 30 min, but that nephrectomy resulted in a significantly reduced clearance, such that detectable serum concentrations of DOM were still present at 60 min postinjection (the last time point studied). In a more robust study of renal pharmacokinetics, Suzuki and Hierlihy (1993) reported that serum clearance of DOM was almost entirely due to renal excretion, with similar kinetics over a wide range of dosages (0.5 ng/kg-2.0 mg/kg i.v.). Analysis of serum concentration-time curves produced elimination rate constants (k) of  $0.025-0.035 \text{ min}^{-1}$  (equivalent to an elimination half-life of approximately 13 min). These authors also reported total body clearance values of 7.75–10.82 ml/min/kg and renal clearance values ranging from 8.80 to 12.20 ml/min/kg. Renal excretion in this study appeared to be mainly by glomerular filtration because kinetics were not altered by the presence of probenecid. Interestingly, these data in rats may help to explain why comparatively few of the humans who ate contaminated shellfish in 1987 presented with symptoms of toxicity (see Sect. 4.2 below).

## 3.2 Acute Behavioral Toxicity

Following the 1987 outbreak of human toxicity following consumption of shellfish contaminated with DOM (see Sect. 4.2 below), a number of studies were published describing the acute toxicity of DOM in mice, rats, and, to a lesser extent, nonhuman primates. A complete review of these studies is beyond the scope of this chapter, but the reader is referred to Doucette and Tasker (2008) and Tasker et al. (2011) for greater detail. One important thing to note when reviewing the early (pre-1995) literature is that it was intended to describe toxicity resulting from contaminated shellfish. Consequently, most of these investigators did not have access to purified DOM, and/or the precise nature of the toxin(s) causing amnesic shellfish poisoning (see Sect. 4.2) was unknown. For these reasons toxicity data are derived largely from intraperitoneal (i.p.) injections of crude homogenates of toxic mussels (Iverson et al. 1989, 1990; Glavin et al. 1989, 1990; Grimmelt et al. 1990; Bose et al. 1990; Tasker et al. 1991; Strain and Tasker 1991) or mussel

hepatopancreas (Bose et al. 1990; Glavin et al. 1990). Variability in sample preparation probably accounts for some of the variability in dose–response data in these chapters. In fact, Novelli et al. (1992) reported that domoic acid toxicity in mussel extracts could be altered by the presence of other amino acids, implying that even identically prepared extracts could have small differences in toxicity if different batches of shellfish are used. This is consistent with detailed dose–response data compiled by Tasker et al. (1991) in which these authors reported differences in the dose–response curves generated using DOM from extracts of toxic mussels (concentrations confirmed by HPLC) and extracts of "clean" mussels that were "spiked" with known concentrations of purified DOM.

Despite the above-mentioned limitations in early studies, subsequently confirmed in later studies using chemically purified DOM, the acute behavioral toxicity elicited by DOM is remarkably consistent both within and between species. Systemic injections of DOM elicit a graded series of behavioral changes prior to seizure onset whether studied in mice (Iverson et al. 1989; Tasker et al. 1991), rats (Tryphonas and Iverson 1990; Scallet et al. 2004), or nonhuman primates (Tryphonas et al. 1990a, b; Scallet et al. 1993; Schmued et al. 1995; Slikker et al. 1998). The sequence of events is both highly reproducible and predictably doserelated. Following injection, animals will initially demonstrate hypoactivity and sedation (accompanied by gagging and vomiting in nonhuman primates, but not seen in rodents) followed by akinesia, rigidity, stereotypies (repetitive scapular scratching in rodents), loss of postural control, convulsions (forelimb only, progressing to full tonic-clonic), and death. Both the extent to which the animal progresses through this sequence and the speed of progression are dose dependent (Tasker et al. 1991). Further, the profile of acute behavioral toxicity does not appear to be appreciably different in male or female animals, although it has been suggested that different strains of mice may demonstrate altered sensitivity to DOM (Peng et al. 1997).

Predictably, behavioral toxicity following acute injections of DOM correlates with abnormal hippocampal discharges and limbic seizures as recorded electroencephalographically (Nakajima and Potvin 1992; Fujita et al. 1996; Scallet et al. 2004). Moreover, and consistent with the effects observed in human intoxication (see Sect. 4.2), acute injections of DOM in rats have also been reported to alter neurobehavioral performance in tests of passive avoidance, auditory startle, or conditioned avoidance response (Sobotka et al. 1996) or of learning and memory (Sutherland et al. 1990; Petrie et al. 1992; Nakajima and Potvin 1992; Clayton et al. 1999).

## 3.3 Histopathology

Despite the differences in study protocol, there is an overall agreement regarding the histopathology of the acute brain lesions associated with DOM toxicity. DOM induces neurodegenerative changes consisting of neuronal shrinkage, vacuolization of the cytoplasm, cell drop out, edema, and microvacuolation of the neuropil. Light and electron microscopic observations in animal studies show two types of neuronal degeneration associated with DOM toxicity: swollen, vacuolated neurons are described intermingled with some shrunken and darkly stained electron dense neurons (Tryphonas et al. 1990a, b; Strain and Tasker 1991; Ananth et al. 2001; Pulido 2008). These findings are consistent with the view that the neuronal degeneration induced by DOM is mostly necrotic, at least in the short term. Studies using markers of cell injury and apoptosis also support this view (Ananth et al. 2001) as do studies in organotypic hippocampal slice cultures (see Sect. 2 above). These changes have preferential distribution within structures of the limbic system (Tryphonas et al. 1990a, b; Tryphonas and Iverson 1990; Strain and Tasker, 1991), and the hippocampus among other brain regions appears to be a specific target site having high sensitivity to DOM toxicity, particularly the pyramidal neurons in the CA3 region, followed by the dentate gyrus and CA1 region (Tryphonas, et al. 1990a, b; Sutherland et al. 1990; Strain and Tasker 1991). With some exception, the CA2 region is reported as the least affected (Slikker et al. 1998). Other regions affected include the piriform and entorhinal cortices, olfactory bulbs, nucleus accumbens, arcuate nucleus, area postrema, and the retina (Iverson et al. 1989; Tryphonas et al. 1990a, b; Tryphonas and Iverson 1990; Schmued et al. 1995). The anatomical extent of brain lesions induced by DOM has also been identified by magnetic resonance imaging microscopy in both human (Cendes et al. 1995) and rats (Lester et al. 1999) and is consistent with the histopathology data.

In addition to neuronal degeneration, there is also evidence suggesting involvement and injury of glial cells, including astrocytes and microglia (Mayer 2000; Mayer et al. 2001; Gill et al. 2008; Pulido 2008) which is also consistent with results obtained in organotypic slice cultures (see Sect. 2 above).

## 3.4 DOM as a Developmental Neurotoxin

Although much of the early DOM literature dealt with acute, and to a lesser extent chronic, toxicity in adult animals (see Sects. 3.2 and 3.3 above), evidence and interest in the effects of DOM on the developing nervous system and DOM's potential role as a developmental neurotoxin have been appearing at an escalating pace over the past several years (for review see Costa et al. 2010). Despite an early report of negative consequences to DOM exposure in utero (Dakshinamurti et al. 1993), the issue of gestational toxicity lays somewhat dormant for many years but has recently been explored in greater detail with publications on maternal-fetal transfer rates (Maucher and Ramsdell 2007) and fetal kinetics (Maucher-Fuquay et al. 2012) as well as reports of toxicity following exposure in utero in rats (Tanemura et al. 2009) and sea lions (Ramsdell and Zabka 2008).

Because rat brain development is delayed relative to humans, early postnatal life in the rat corresponds roughly to late third-trimester brain development in humans, a particularly sensitive period of brain development. Pioneering studies by Xi et al. (1997) and Doucette et al. (2000) demonstrated convincingly that the neonatal rat brain was exquisitely sensitive to the toxic effects of DOM. Further, this difference in potency relative to adult rats changes throughout early life, with dose-response relationships for DOM toxicity shifting progressively to the right between 0 and 22 days of age (Doucette et al. 2000), an effect that is presumed to be due to maturation of the blood-brain barrier. During the second postnatal week of life in the rat (i.e., the middle of the brain growth spurt; Dobbing and Smart 1974), doses of DOM as low as 20 µg/kg can alter neurobehavioral development (Doucette et al. 2003). Of particular interest, and concern, are an ever-increasing number of publications showing that exposure of rat pups to low doses of DOM during early development can result in later-onset symptoms of disease in the resulting adult animal. Best characterized to date are the effects of low-dose neonatal DOM in producing a neurodevelopmental model of temporal lobe epilepsy characterized by changes in behavior (Doucette et al. 2004), hippocampal morphology (Doucette et al. 2004; Bernard et al. 2007; Gill et al. 2010b), seizure threshold (Gill et al. 2010a), and cognition (Gill et al. 2012). More recently, low doses of DOM during development have also been shown to produce apparently permanent changes in both sleep patterns (Gill et al. 2009) and attentional processing performance (e.g., prepulse inhibition, latent inhibition) (Adams et al. 2008; Marriott et al. 2012) that are commonly seen in schizophrenia and related neuropsychiatric disorders. These observations and the resulting slowly developing disease models can prove useful in dissecting the complex sequence of changes that underlies neurological diseases, but given that current regulatory limits on DOM are based on acute toxicity in adult animals, they also raise serious questions about the potential of low concentrations of DOM in the environment or in food being a developmental neurotoxin that may warrant additional scrutiny (Stewart 2010).

## 4 Clinical Toxicity

## 4.1 Toxicity in Wildlife

DOM and to some extent DOM isomers have been found in oceans worldwide, so it is not surprising that there have been many reports of DOM toxicity in wildlife species. Reviews of much of this data can be found in Doucette and Tasker (2008) and Lefebvre and Robertson (2010) as well as other sources. While production of DOM in the aquatic environment has historically been associated with the red alga *Chondria armata* (Takemoto and Daigo 1960), the source of DOM as an environmental toxicant is through blooms of planktonic species including *Pseudo-nitzschia multiseries* and *Pseudo-nitzschia australis* as well as many others. For further detail, the reader is directed to any of several excellent reviews including Bates and Horner (1998) and Bates and Trainer (2006). Plankton species are a food source for many aquatic invertebrates particularly bivalve molluscs (e.g., mussels, clams, oysters) as well as planktivorous fish (e.g., anchovies, sardines) and mammals (e.g., filterfeeding whales). While toxicity has been reported in all of these species, it is also clear that DOM toxicity extends higher up to the food chain and affects invertebrates that consume contaminated shellfish (e.g., crabs) as well as piscivorous birds (e.g., cormorants, pelicans) and marine mammals that eat fish and/or shellfish (e.g., sea lions, sea otters, piscivorous whales) (Lefebvre et al. 1999, 2000, 2001, 2002).

The symptoms associated with DOM toxicity in these species vary, but it is generally believed that most invertebrate species are unaffected (presumably due to their rudimentary nervous system), whereas fish, birds, and mammals display classic signs of neurotoxicity. Indeed it is believed, although not confirmed, that seabirds intoxicated with DOM in California were the inspiration for Alfred Hitchcock's famous movie The Birds (<sup>®</sup> Universal Pictures, 1963). Less romantic, but far better documented, accounts of DOM intoxication in wildlife have been reported by a number of groups, including an excellent series of reports on intoxication of California sea lions (*Zalophus californianus*) by both US west coast and east coast scientists (e.g., Scholin et al. 2000; Ch'ng et al. 2002; Gulland et al. 2002; Silvagni et al. 2005; Goldstein et al. 2008).

## 4.2 Toxicity in Humans

Late in 1987, an outbreak of a newly recognized acute illness caused by eating blue mussels and characterized by gastrointestinal and unusual neurological symptoms occurred in Canada. More than 107 people (47 men and 60 women) were affected, most from Quebec. A case was defined as the presence of gastrointestinal symptoms (vomiting, abdominal cramps, diarrhea) within 24 h and neurological symptoms within 48 h (severe headache and memory loss). The etiologic agent was found to be DOM, an excitatory neurotransmitter amino acid, produced by *Nitzschia pungens f. multiseries*. Both *N. pungens* and DOM were present in the digestive glands of the causative cultivated mussels, harvested from the eastern coast of Prince Edward Island, and shipped to other parts of Canada (Perl et al. 1990; Teitelbaum et al. 1990).

Of the 107 persons that met the case definition, 19 were hospitalized from 4 to 101 days, 12 of these were admitted to intensive care, and 3 died in hospital at 12–18 days after admission. Another patient died after 3 months. Ninety nine of the patients answered a questionnaire, in which they provided information on the symptoms and on the amount of mussels consumed. Of these patients, 49 were between 40 and 59 years old and 38 patients were 60 years or older. Symptoms of illness included nausea (77 %), vomiting (76 %), abdominal cramps (51 %), diarrhea (42 %), headache (43 %), and memory loss (25 %). None of the younger patients (20-39 yrs) suffered memory loss, and their only symptoms were of a gastrointestinal nature. Hospital charts were available for 16 of the 19 hospitalized patients, indicating that all severely ill patients less than 65 years old had preexisting illnesses, including diabetes (3), chronic renal disease (2), and hypertension with a history of transient ischemic attacks (1) considered to be predisposing factors. All patients admitted to intensive care had serious neurologic dysfunction, including coma (9), mutism (11), and seizures (8). Seven patients had unstable blood pressure or cardiac arrhythmias. For nine patients and one person who did not become ill, analytical information on the unconsumed portion of the

Patient	Age	Estimated DOM consumed (mg)	Clinical symptoms and treatment			
			GI	Memory loss	Hospitalized	ICU
1	72	60	Y	-	-	-
2	62	70	Y	Y	-	-
3	70	80	Y	-	-	-
4	61	90	Y	-	-	-
5	67	110	Y	_	-	-
6	71	110	Y	-	-	-
7	74	270	Y	Y	Y	-
8	68	290	Y	Y	Y	Y
9	84	290	Y	Y	Y	Y

**Table 1** Estimated exposure and clinical course of patients who ingested DOM in contaminated mussels during the 1987 outbreak of amnesic shellfish poisoning in Canada

Data adapted from Perl et al. (1990)

GI refers to gastrointestinal symptoms; ICU refers to intensive care unit

mussels and recall information on portion size were available, and this was used to estimate exposure. The concentration of DOM in these mussels was determined by mouse bioassay (characteristic hind leg scratching) and ranged between 31 and 128 mg/100 g. Increasing exposure correlated with the clinical course of events (Table 1). All patients reported gastrointestinal illness, but only one of six patients who consumed between 60 and 110 mg DOM suffered memory loss and none required hospitalization. All three patients who had consumed 270–290 mg DOM suffered neurological symptoms and were hospitalized. One person who consumed only 20 mg DOM did not become ill. The cognitive impairment observed in this new disease, attributed to DOM, appeared to be persistent and led to the term "amnesic shellfish poisoning" (Perl et al. 1990; Todd 1990). For a detailed description of neurologic and neuropsychiatric profiles of patients subsequent to the initial event, the reader is referred to reports by Gjedde and Evens (1990) and Cendes et al. (1995).

Neuropathological examination of the four patients who died indicated neuronal necrosis and astrocytosis particularly in the hippocampus and the amygdaloid nucleus. All four victims also had lesions in the claustrum, secondary olfactory areas, the septal area, and the nucleus accumbens septi. Two had prominent thalamic damage, especially in the dorsal medial nucleus. The subfrontal cortex was also damaged in three of the patients. The authors noted that the pattern of damage in the hippocampus appeared to parallel that seen in animals which suffered neurotoxic reactions after administration of kainic acid. Cerebrovascular disease was evident in two patients: an acute hemorrhagic infarct in the left posterior perisylvian cerebral in one victim and, in the other, cerebellar infarcts likely related to a myocardial infarct 3 days prior to death. Neurofibrillary tangles, and senile plaques, typical features of Alzheimer's disease, were not observed in these patients (Teitelbaum et al. 1990).

Although the 1987 incident in Montreal is the only confirmed case of human intoxication with DOM, an outbreak of DOM poisoning may have occurred in

October/November 1991 in Washington State, USA. Approximately two-dozen people became ill after ingesting razor clams harvested along the Washington and Oregon coasts. Although gastrointestinal or neurological symptoms were observed within 36 h of ingestion, no other symptoms consistent with DOM intoxication were reported. A total of 21 possible DOM victims and 43 non-victims were interviewed. Based on the concentrations of DOM measured in razor clams, the mean total DOM consumption by the 21 victims would have been 17 mg (estimated to be 0.28 mg/kg b.w.) and 8 mg by the unaffected individuals (0.13 mg/kg b.w.). A total of 13 patients developed neurological symptoms, but all recovered. Unfortunately the incident was not well reported and was unsubstantiated (Todd 1993; Jeffrey et. al. 2004).

## 5 Conclusion

Revived from relative obscurity into the limelight as a result of the 1987 incident of human intoxication, DOM has become an environmental and experimental toxin of considerable importance worldwide. As an environmental contaminant affecting both the food supply and wildlife species, DOM and DOM isomers are now routinely monitored in many jurisdictions. As an experimental tool in neurotoxicology and neuroscience, DOM has a unique place as a slowly desensitizing AMPA/KA-R agonist that shows high potency and efficacy in a wide variety of experimental systems. Perhaps the most interesting thing, and of great potential concern, however, is the emerging evidence of DOM as a neurodevelopmental toxin and a precipitating factor in the onset and progression of various forms of neurological disease. Whether these properties can be exploited in the form of disease models and/or warrant improved vigilance on the part of global regulatory bodies, continued study of the many fascinating properties of DOM and isomers is warranted.

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# Endogenous Kynurenic Acid and Neurotoxicity

## Ewa M. Urbańska, Iwona Chmiel-Perzyńska, Adam Perzyński, Marek Derkacz, and Björn Owe-Larsson

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E.M. Urbańska (⊠)

Department of Toxicology, Institute of Agricultural Medicine, Lublin, Poland e-mail: emurbanska@poczta.onet.pl

I. Chmiel-Perzyńska

Laboratory of Cellular and Molecular Pharmacology, Department of Experimental and Clinical Pharmacology, Medical University of Lublin, Lublin, Poland e-mail: iwona.perz@poczta.onet.pl

#### A. Perzyński Department of Psychiatry, Medical University of Lublin, Lublin, Poland e-mail: adam.perzynski@wp.pl

M. Derkacz Department of Endocrinology, Medical University of Lublin, Lublin, Poland e-mail: donmarcos@wp.pl

B. Owe-Larsson

Department of Clinical Neuroscience, Karolinska Institutet, Section of Psychiatry at Karolinska University Hospital Huddinge, Stockholm, Sweden e-mail: bjorn.owe-larsson@ki.se

Laboratory of Cellular and Molecular Pharmacology, Department of Experimental and Clinical Pharmacology, Medical University of Lublin, Lublin, Poland

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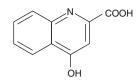
### Abstract

Tryptophan metabolism along kynurenine pathway yields a number of compounds affecting brain function. Among kynurenine derivatives, neuroprotective kynurenic acid (KYNA) and neurotoxic quinolinic acid and 3-hvdroxykynurenine have stimulated the greatest scientific interest. KYNA, initially considered merely a side-product of tryptophan degradation, was discovered in 1982 to act as excitatory amino acid receptor antagonist. Since then, a number of novel KYNA targets emerged. KYNA was suggested to play a role as antagonist of  $\alpha$ 7 nicotinic receptors and ligand of G protein-coupled GPR35 and human aryl hydrocarbon (AHR) receptors. In here, research data is reviewed supporting the idea that produced by astrocytes KYNA serves as an endogenous neuroprotectant. Mechanisms controlling brain levels of KYNA are discussed in the context of neurodegenerative disorders, brain ischemia, and seizures. Available data concerning changes of brain KYNA in respective animal models and in human diseases, together with an overview of effects following the application of KYNA, KYNA analogues or compounds influencing the activity of enzymes along kynurenine pathway are presented. Emerging therapies designed to increase the level of neuroprotective KYNA may become an important avenue in the treatment of brain disorders accompanied by neuronal loss.

## 1 Introduction

The history of kynurenic acid (KYNA; see Fig. 1) dates to year 1853, when German chemist Baron Justus von Liebig determined its presence in canine urine (Liebig 1853). Half a century later, the discovery of tryptophan was followed by the observation that KYNA is one of its metabolites (Hopkins and Cole 1901; Ellinger 1904). For almost eight decades, the compound was considered merely a side-product of *tryptophan* degradation. In early 1980s, KYNA was reported to antagonize amino acid-induced excitation of central neurons (Perkins and Stone 1982) and to block neurotoxicity and seizures evoked by another tryptophan metabolite, quinolinic acid (Foster et al. 1984). Soon, the presence of KYNA within the brain of mammals and rodents was confirmed (Moroni et al. 1988a, b; Turski et al. 1988). Subsequently, numerous basic and clinical investigations addressed the role of KYNA in brain function, and the potential ways to modulate KYNA central levels were elucidated.

Fig. 1 Structure of kynurenic acid



## 2 KYNA Synthesis and Brain Levels

Tryptophan, after entering the brain in a free form, is utilized along various metabolic pathways yielding structural and enzymatic proteins or centrally active substances such as serotonin and kynurenines. In mammals, approx. 95–99 % of dietary tryptophan not bound into proteins is converted to kynurenine along the so-called *kynurenine* pathway (Stone and Darlington 2002); see Fig. 2.

Brain kynurenine derives from production in situ (40–50 %) and from peripheral pool. Kynurenine enters the brain via large neutral amino acid transporter (Fukui et al. 1991; Németh et al. 2006). KYNA itself penetrates across the blood-brain barrier to a lesser degree (Scharfman and Goodman 1998; Salvati et al. 1999). However, peripheral application of KYNA prior to ischemia may dramatically increase brain concentrations of compound (Salvati et al. 1999). Transport occurs mostly via passive diffusion (Fukui et al. 1991). Notably, human organic anion transporter hOT3 might play a role in the uptake of KYNA from the brain into brain capillary endothelial cells (Uwai et al. 2012).

Within the brain, kynurenine can be either transaminated to KYNA or hydroxylated to *3-hydroxykynurenine*, which is further converted to *quinolinic acid*, and finally to nicotinamide adenine dinucleotide (NAD) (Moroni 1999; Németh et al. 2005, 2006). 3-Hydroxykynurenine generates large quantities of free radicals, whereas quinolinic acid is an *N*-methyl-D-aspartate receptor agonist with potent excitotoxic properties (Stone 2001). Of many kynurenine derivatives, neuroinhibitory KYNA and neurotoxic quinolinic acid and 3-hydroxykynurenine have stimulated the greatest scientific interest.

The brain level of KYNA depends directly on the activity of kynurenine aminotransferases (KATs), but, indirectly, it is also strongly associated with the action of enzymes contributing to the production and degradation of its precursor, kynurenine. The synthesis of kynurenine is regulated by two separate enzymes, tryptophan 2,3-dioxygenase (TDO) and indoleamine 2,3-dioxygenase (IDO). TDO and IDO are expressed in the brain but their activities are rather low. Therefore, as mentioned above, more than 50 % of brain kynurenine is regarded to originate from peripheral sources.

Increasing scientific interest has focused recently on *kynurenine* 3-monooxygenase (KMO), which converts kynurenine to 3-hydroxykynurenine. As KMO displays lower  $K_m$  for kynurenine than KATs (Bender and McCreanor 1985), it is considered a key enzyme controlling the availability of kynurenine for KYNA synthesis. The process of irreversible transamination of kynurenine to

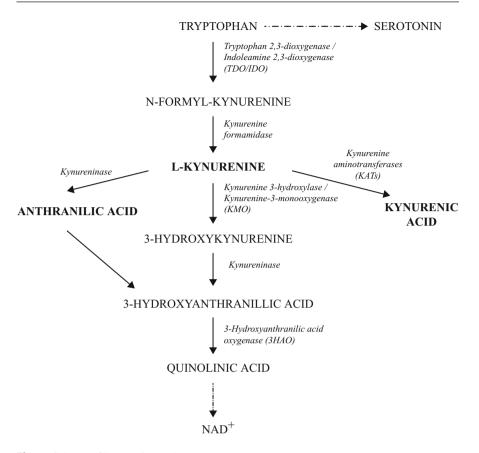


Fig. 2 Scheme of kynurenine pathway

KYNA is catalyzed by four distinct kynurenine aminotransferases I–IV (KATs) (Okuno et al. 1991a; Guidetti et al. 2007; Bellocchi et al. 2009) considered nowadays as potential targets for regulating brain KYNA levels. KAT I [glutamine transaminase K (GTK)/cysteine conjugate beta-lyase (CCBL) 1], KAT II [aminoadipate aminotransferase (AADAT)], KAT III (CCBL2), and KAT IV [glutamic-oxaloacetic transaminase (GOT) 2/mitochondrial aspartate amino-transferase (mASPAT)] have been identified in mouse, rat, and human brains (Han et al. 2010). KATs are expressed primarily in the *astrocytes* (Roberts et al. 1992; Guidetti et al. 1997), but neurons are also able to produce KYNA (Du et al. 1992; Roberts et al. 1992; Rzeski et al. 2005). KATs display diverse optimal pH range and different levels of specific activity in specific brain regions and, although sharing a number of amino acid and  $\alpha$ -ketoacid substrates, have distinct substrate profile (Okuno et al. 1991b; Guidetti et al. 2007; Bellocchi et al. 2009).

Concentrations of KYNA in the central nervous system (CNS) vary depending on the analyzed species, age, and CNS area. In brain structures, KYNA levels range from 0.014 to 3.38 pmol/mg of tissue (Carlá et al. 1988; Turski et al. 1988; Swartz et al. 1990; Beal et al. 1992; Jauch et al. 1995). The lowest concentrations of KYNA in the human brain were detected in cerebellum and medulla (approx. 0.1–0.3 pmol/mg). In cortical areas and substantia nigra, KYNA reaches approx. 0.2–0.6 pmol/mg, whereas the highest concentrations occur in globus pallidus and putamen (0.7–1.4 pmol/mg) (Turski et al. 1988; Swartz et al. 1990; Jauch et al. 1995; Baran et al. 1996). The content of KYNA in *cerebrospinal fluid* (CSF) ranges from 1 to 5 nM (Swartz et al. 1990; Erhardt et al. 2001; Rejdak et al. 2002; Hżecka et al. 2003).

The level of KYNA in the CNS increases with age. In humans aged <50 years, KYNA content in CSF is lower in comparison with elderly individuals (2.8 vs. 4.1 fmol/µl) (Kepplinger et al. 2005). In rats between 1st week and 18th month of age, KYNA increases over 50-fold, from approx. 15 to 750 pmol/g of protein (Moroni et al. 1988a). Others reported rise of brain KYNA by 80 % up to 350 %, between 3rd and 24th month of age, from approx. 20 to approx. 70 pmol/g of tissue (Gramsbergen et al. 1992; Braidy et al. 2011).

In periphery, the presence of KYNA was demonstrated in various animal and human tissues such as kidneys, liver, or heart (Carlá et al. 1988). Large quantities of KYNA are released by vascular *endothelium*, as demonstrated in rat aortic rings and endothelial cell cultures (Stazka et al. 2002; 2005). In humans, serum levels of KYNA range between 4 and 30 nM (Swan et al. 1983; Urbanska et al. 2006; Hartai et al. 2007; Baran et al. 2010). In rats, the concentrations are slightly higher and reach 20–90 nM (Stazka et al. 2005; Fukushima et al. 2009; Baran et al. 2010).

Interestingly, KYNA is not only synthesized de novo by mammalian tissues but also produced by saprophytic bacteria in the intestinal tract and exogenously delivered with food (Kuc et al. 2008; Turski et al. 2009). High concentration of KYNA was found in food, e.g., in herbs, bee products, broccoli, and potatoes (Turski et al. 2009).

#### **3** Regulation of KYNA Synthesis

Brain level of KYNA may be regulated in various ways, including the availability of direct precursor and the activity of biosynthetic enzymes. Accumulating data identified also a number of other modulatory factors, of either endo- or exogenous origin.

Extracellular milieu, including ionic composition, pH, glucose, and oxygen availability, can affect KYNA production, as shown in vitro in brain cortical slices (Turski et al. 1989; Gramsbergen et al. 1991, 1997). Low concentrations of Cl<sup>-</sup> and K<sup>+</sup> as well as depolarizing concentrations of K<sup>+</sup> (50 mM) reduce, whereas high Na<sup>+</sup> increases KYNA production. Hypoxia, anoxia, and hypoglycemia inhibit KYNA synthesis, while hyperglycemia remains without effect (Turski et al. 1989; Chmiel-Perzyńska et al. 2007). It has also been shown that  $\beta$ -hydroxybutyrate, one of the ketone bodies rising in the serum during diabetic ketoacidosis or *ketogenic diet*, stimulates brain KYNA synthesis in cortical slices and glial cultures (Chmiel-Perzyńska et al. 2011).  $\beta$ -hydroxybutyrate did not affect directly the activity of semi-purified preparation of KYNA biosynthetic enzymes in vitro. However, in

glial cultures  $\beta$ -hydroxybutyrate stimulated the de novo expression of KAT I and II. These data suggest that during fasting or ketogenic diet, high level of  $\beta$ -hydroxybutyrate may not only protect brain from the reduction of KYNA elicited by hypoglycemia and/or acidosis but actually enhance its synthesis above control values (Chmiel-Perzyńska et al. 2011).

Endogenous agonists of ionotropic glutamate receptors, such as *L-cysteine* sulfinate, L-aspartate, L-glutamate, L-cysteate, L-homocysteine sulfinate, and L-homocysteate, reduce the production of KYNA in cortical slices with IC<sub>50</sub> values of 0.08, 0.3, 0.79, 1.77, 2.9, and 3.93 mM, respectively (Urbanska et al. 1997, 2000; Kocki et al. 2003). Their action is mediated most probably via inhibition of KAT I/ KAT II activity. Among studied agonists, L-cysteine sulfinate appears one of the most potent KAT II inhibitors, displaying the IC<sub>50</sub>=2  $\mu$ M (Kocki et al. 2003).

Synthetic *metabotropic receptor agonists* were also shown to reduce KYNA synthesis in cortical slices. Among them, quisqualate was unequivocally the most potent inhibitor, acting in the micromolar range of concentrations, while agonists of group I (DHPG), group I/II (t-ACPD), group II (L-CCG-I), and group III (L-AP4) acted weaker (Urbanska et al. 1997). Interestingly, L-AP4 and t-ACPD were ineffective in spinal cord slices (Urbanska et al. 2000).

The effect of *dopaminergic* ligands on striatal KYNA production was studied in vivo in developing and adult rats (Rassoulpour et al. 1998). D<sub>1</sub> receptor antagonist SCH 23390, but not the D<sub>2</sub> receptor antagonists, raclopride and haloperidol, increased striatal KYNA levels, whereas D<sub>1</sub> and D<sub>2</sub> receptor agonists reduced KYNA levels approximately to the same extent in young animals. D-Amphetamine impaired striatal synthesis of KYNA irrespectively of animals' age (Rassoulpour et al. 1998). Similarly, peripheral application of L-DOPA, precursor of dopamine, decreased extracellular KYNA in rat striatum (Wu et al. 2002). In humans, however, L-DOPA treatment does not seem to affect central KYNA levels (Ogawa et al. 1992; Samadi et al. 2005).

Selective and mixed  $\beta_1$ - and  $\beta_2$ -adrenergic agonists, such as xamoterol, denopamine, albuterol, clenbuterol, or epinephrine, as well as cAMP analogues, enhanced KYNA synthesis in vitro and in vivo (Luchowska et al. 2008, 2009). KT5720, a selective inhibitor of *protein kinase A* (PKA), strongly reduced KYNA formation in cortical slices and in glial cultures. Notably, the effective concentrations were extremely low, ranging between 0.1 and 10 µM. β-adrenergic agonists, cAMP analogues, or KT5720 did not affect directly the activity of KATs measured in vitro, in semi-purified homogenate. However, the exposure of cultured glial cells to cAMP analogues and β-adrenergic agonists stimulated the enzymatic conversion of kynurenine, most probably due to an enhancement of KATs synthesis. The cAMP/PKA-related pathway seems an important factor regulating the endogenous levels of KYNA via expression of KATs (Luchowska et al. 2008, 2009).

*Nitric oxide* (NO) donors, S-nitroso-N-acetylpenicillamine (SNAP), and 3morpholinosydnonimine (SIN-1) enhanced KYNA production in cortical slices yet did not affect KATs activity (Luchowski and Urbanska 2007). SNAP is the source of NO and S-nitrosylating agent, whereas SIN-1 is the NO/peroxynitrite (ONOO–) generator with a very weak S-nitrosylating activity. Their stimulatory effects were abolished in the presence of free radicals scavenger, L-ascorbate, indicating possible involvement of the excessive release of reactive oxygen species in the observed phenomenon (Luchowski and Urbanska 2007).

*Mitochondrial toxins* compromising the status of mitochondrial respiration, such as aminooxyacetic acid (AOAA), *3-nitropropionic acid (3-NPA)*, or *1-methyl-4phenylpyridinium (MPP<sup>+</sup>)*, inhibit KYNA synthesis in vitro and in vivo (Turski et al 1989; Urbanska et al. 1997; Luchowski et al. 2002; Kocki et al. 2003). AOAA disrupts the function of aspartate-malate shuttle across the mitochondrial membrane; 3-NPA is an irreversible inhibitor of succinate dehydrogenase and mitochondrial complex II, whereas MPP<sup>+</sup> blocks the activity of complex I and IV (Urbanska 2005). 3-NPA and AOAA inhibit the activity of KAT I and KAT II, whereas MPP<sup>+</sup> blocks only KAT II action (Luchowski et al. 2002). Interestingly, mitochondrial toxins exert indirect excitotoxic effects, modeling Parkinson's (MPP<sup>+</sup>) and Huntington's (AOAA and 3-NPA) diseases upon local or intracerebral application (Urbanska 2005).

*D,L-homocysteine*, which increased levels are considered a risk factor in atherosclerosis, biphasically affects KYNA production in the endothelium. In micromolar concentrations (approx. 40–100  $\mu$ M), D,L-homocysteine stimulates, whereas in higher concentrations, it inhibits KYNA formation (Stazka et al. 2005). Similarly, low concentrations of compound stimulated KYNA synthesis, while millimolar concentrations impaired KYNA formation (IC<sub>50</sub>= approx. 5–6 mM) in the brain (Luchowska et al. 2005; Chmiel-Perzyńska et al. 2007). Metabolic derivative of D,L-homocysteine, *S*-adenosylhomocysteine, displayed similar but more potent activity than parent compound, augmenting KYNA production at 0.03–0.08 mM and reducing it at  $\geq$ 0.5 mM. The stimulatory effect of *S*-adenosylhomocysteine was abolished in the presence of the kynurenine uptake inhibitors, L-leucine and L-phenyloalanine (Luchowska et al. 2005).

Number of clinically effective compounds may exert their therapeutic effects, at least in part, via increase of central KYNA synthesis. *Memantine*, a well-tolerated derivative of amantadine acting predominantly as an open-channel blocker of *N*-methyl-D-aspartate (NMDA) receptors, protects neurons against excitotoxic insults under different experimental paradigms in vitro and in vivo (Danysz and Parsons 2003). The compound is used clinically to ameliorate symptoms of Parkinson's and Alzheimer's diseases. Recently, memantine was found to stimulate KYNA production at low, micromolar concentrations, possibly via PKA-dependent mechanism (Kloc et al. 2008).

Antiepileptic drugs, carbamazepine, felbamate, phenobarbital, phenytoin, and lamotrigine (at 0.5–3.0 mM), were found to increase KYNA synthesis in rat cortical slices and to enhance KAT I activity (Kocki et al. 2004, 2006). On the contrary, vigabatrin, gabapentin, and tiagabine decreased KYNA production, possibly due to the inhibition of KAT I and KAT II. This aspect of their action may be partially involved in the paradoxical exacerbation of absence or myoclonic seizures, as sometimes observed clinically after administration of vigabatrin, gabapentin, and tiagabine (Kocki et al. 2006).

Recently, *antidepressants* fluoxetine, citalopram, amitriptyline, and imipramine  $(1-10 \ \mu\text{M})$  were demonstrated to increase de novo production of KYNA and

to diminish 3-hydroxykynurenie synthesis (at 24 and 48, but not 2-h incubation). RT-PCR studies revealed that antidepressants did not alter KAT1, KAT2 and kynurenine-3-monooxygenase (Kmo) gene expressions after 2-h exposure. After 48 h, the expression of KAT1 and KAT2 was strongly upregulated, and Kmo expression was downregulated by all antidepressants. These findings indicate that antidepressants may act in part via restoring the beneficial ratio between KYNA and 3-hydroxykynurenine (Kocki et al. 2012).

Notably, also the environmental factors may affect central KYNA production. Pyrethroid *pesticides*, such as deltamethrin and fenpropathrin, were shown to inhibit KYNA production in cortical slices (Zielińska et al. 2005). Tobacco smoking may either increase or decrease central KYNA levels, depending on the exposure time. In rats, short, 4- to 6-day administration of nicotine decreased brain KYNA. Withdrawal of nicotine resulted in the return of KYNA concentration to control values (Rassoulpour et al. 2005). In contrast, 10-day exposure to nicotine increased KYNA level in hippocampus, striatum, and cortex, but not in the serum (Rassoulpour et al. 2005). Others confirmed the stimulatory effect of 10-day nicotine application. However, chronic, 30-day exposure to nicotine was reported to increase serum KYNA level and to decrease cortical KYNA content, despite the enhanced KATs activity (Zielińska et al. 2009). Only central, but not the plasma changes of KYNA, were reversed by the use of an antagonist of nicotinic receptors, mecamylamine (Zielińska et al. 2009). Possibly, nicotine application induces distinct mechanisms involved in the regulation of KYNA synthesis, distribution, and elimination (Zielińska et al. 2009).

### 4 Mechanisms of KYNA Action

Since the initial discovery identifying KYNA as en endogenous antagonist of ionotropic glutamate receptors (Perkins and Stone 1982), a number of novel KYNA targets emerged. Currently, KYNA is also suggested to act as an antagonist of nicotinic receptors and as ligand of G protein-coupled GPR35 and human aryl hydrocarbon (AHR) receptors.

In electrophysiological studies, KYNA was shown to block competitively the activation of ionotropic glutamate receptors of *NMDA*,  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (*AMPA*) and kainate type, at relatively high  $\mu$ M concentrations (Perkins and Stone 1982, 1983; Stone 1993). At much lower concentrations (10–30  $\mu$ M; IC<sub>50</sub>=8  $\mu$ M), KYNA acts as an antagonist of the strychnine-insensitive glycine site of NMDA receptor complex (Kessler et al. 1989; Stone 1993). Bearing in mind that extracellular concentration of KYNA is within nM range, initially endogenous KYNA levels were considered too low to modulate excitatory neurotransmission. However, accumulated data leave no doubt that even modest increase of KYNA may alter brain function, e.g., change the neuronal firing in locus coeruleus, reduce the locomotor activity, impair cognition, and prevent seizures or impair excitotoxic neuronal loss (Vécsei and Beal 1991; Russi et al. 1992; Carpenedo et al. 1994, 2002; Pocivavsek et al. 2011).

Moreover, locally applied low nanomolar concentrations of KYNA were shown to reduce the release of glutamate in caudate nucleus and to inhibit the neurotransmitter release in striatal preparations (Carpenedo et al. 2001). Experimental studies confirmed that fluctuations of KYNA level may alter glutamine, acetylcholine, and dopamine release (Wonodi and Schwarcz 2010).

KYNA was also reported to antagonize the  $\alpha_7$  *nicotinic receptors* (nAChRs) in hippocampus, already at low 100 nM concentration, reaching IC<sub>50</sub>~7 µM. However, KYNA effect in slices was much less robust than that seen in cultured neurons (Hilmas et al. 2001; Stone 2007). Hippocampal  $\alpha_7$  nAChRs are located on some GABAergic neuronal cell bodies and on glutamatergic dendrites and axon terminals and seem to modulate neurotransmitter release rather than to mediate neurotransmission (Vizi and Lendvai 1999; Dajas-Bailador et al. 2000). Axonal  $\alpha_7$  nAChRs are involved in the modulation of presynaptic NMDAR expression and structural plasticity of glutamatergic presynaptic boutons during early synaptic development (Lin et al. 2010). The role of KYNA in the modulation of  $\alpha_7$  nAChRs in vivo is yet not clear, because three groups failed to detect any effect of KYNA on  $\alpha_7$  nAChRsmediated events using electrophysiological methods and direct patch-clamp recording in hippocampal cultures and rodent acute brain slices (Arnaiz-Cot et al. 2008; Mok et al. 2009; Dobelis et al. 2012).

In 2006, KYNA was discovered to act as a ligand of G protein-bound receptor, GPR35 (Wang et al. 2006). The possible role of altered GPR35 function in the development of hypertension, immune modulation, and gastric cancer has been suggested (Okumura et al. 2004; Min et al. 2010). GPR35 is highly expressed in human monocytes, T cells, neutrophils, dendritic cells, and within the gastrointestinal tract (Wang et al. 2006). The potency of KYNA toward GPR35 varies depending on experimental paradigm and effector system. In Chinese hamster ovary cells transfected with GPR35, KYNA elevated intracellular Ca<sup>2+</sup> with the  $EC_{50}$  of approx. 7  $\mu$ M (rat) and 40  $\mu$ M (human) (Wang et al. 2006). Using GPR35- $\beta$ -arrestin-2 interaction assay, KYNA was demonstrated to promote the interaction of GPR35 with  $\beta$ -arrestin-2 and cause internalization of the receptor systems with low potency, at  $>100 \ \mu M$  (Jenkins et al. 2011). Others, however, have shown that the arrest of human monocytes on human umbilical vein endothelial cells expressing ICAM-1 (intercellular adhesion molecule 1), in the process mediated by  $\beta_2$ -integrins, occurs at low, 300 nM concentration of KYNA (Barth et al. 2009). Thus, stimulation of GPR35 with KYNA may contribute to the development of inflammatory changes. GPR35 presence was also confirmed within CNS, in the spinal cord and dorsal root ganglia, cerebellum, and brain (Wang et al. 2006; Guo et al. 2008; Ohshiro et al. 2008). In mouse glial cell cultures, the expression of GPR35 receptors was confined to the cytosol and the membranes (Cosi et al. 2011). In these cells, KYNA significantly reduced the forskolin-induced cAMP accumulation with EC<sub>50</sub> of approx. 1 µM (Cosi et al. 2011). Other effects assigned to the activation of GPR35 include voltage-dependent Ca2+-current inhibition (10 µM-1 mM), antinociception, and regulation of neuronal firing (Guo et al. 2008; Zhao et al. 2010). Thus, to date the majority of reports suggest that the interaction of KYNA with GPR35 receptors occurs at concentrations exceeding its extracellular levels. Further research is needed to clarify whether KYNA indeed plays a physiologically important role as a ligand of GPR35.

The latest discovery indicates that KYNA and kynurenine acts as the agonists of *human aryl hydrocarbon receptor (AHR)* (DiNatale et al. 2010; Opitz et al. 2011). AHR is essential for the toxicity of dioxins and related compounds. Additionally, it plays a role in adaptive metabolism and development, hormonal signaling, hematopoiesis, insulin release, and tumor development. Hyperactivation of AHR may lead to a variety of transcriptional responses (Bradshaw and Bell 2009; Bock and Köhle 2009; Stevens et al. 2009; Esser 2012). Stimulation of AHR by exogenous ligands coupled with inflammatory signals can evoke synergistic induction of *interleukin-6* (IL-6) expressions in tumor cells (DiNatale et al. 2010). KYNA was shown to enhance the expression of IL-6 at low 100 nM concentration and was suggested as a factor involved in the escape of tumors, via IL-6-dependent pathway, from the immune surveillance (DiNatale et al. 2010).

Initially considered to act primarily through the inhibition of glutamatemediated neurotransmission, KYNA, with new reports emerging, is not regarded an inert metabolite anymore. Accumulating data support the view that this product of kynurenine pathway may be essentially involved in human physiology and pathology.

## 5 Experimental KYNA Deficiency

In the search for selective tools able to decrease KYNA formation, scientific attention was drawn to aminooxyacetic acid (AOAA) (Urbanska et al. 1998). AOAA, originally recognized as a nonselective inhibitor of aminotransferases (Wood and Peesker 1973), was identified as one of the most potent inhibitors of KYNA synthesis in vitro and in vivo (Turski et al. 1989; Speciale et al. 1990). It blocks KATs activity displaying IC<sub>50</sub> values in very low micromolar range and is only slightly weaker than L-cysteine-sulfinate (Turski et al. 1989; Okuno et al. 1991b; Chmiel-Perzyńska et al. 2007). Using in vivo approach, it was demonstrated that intrastriatally injected AOAA evokes selective excitotoxic quinolinate-like neuronal loss in rats (Urbanska et al. 1989, 1991), what was confirmed by others (Beal et al. 1991). Neurochemical, histological, and behavioral features of striatal neuronal loss closely resembled changes observed in the course of Huntington's disease (HD) (Urbanska et al. 1989, 1991; Beal et al. 1991). Lesions blocked with the use of KYNA and NMDA antagonists did not occur following ablation of corticostriatal glutamatergic input and were age dependent (Urbanska et al. 1989, 1991). Bilateral lesions of caudate with AOAA markedly affected cataleptic response of animals, increasing the reaction to arecoline and morphine and reducing the response to haloperidol (Urbanska et al. 1989, 1991), mimicking the behavioral changes characteristic for other HD models (Sanberg 1980; Sanberg et al. 1981). Bearing in mind that AOAA is a nonspecific inhibitor of aminotransferases, AOAA-induced effects could be attributed to the inhibition of enzymes other than KATs. However, neither application of GABA aminotransferase inhibitor nor L-aspartate aminotransferase inhibitor evoked neuronal degeneration in striatum (Urbanska et al. 1989, 1991). It was also proposed that AOAA-induced indirect excitotoxicity results from mitochondrial dysfunction, as AOAA is known to disrupt the malate-aspartate shuttle across mitochondrial membrane (Beal et al. 1991). However, AOAA affects KYNA synthesis at much lower concentration than required to alter mitochondrial respiration (Hotta 1968). Thus, the diminished formation of KYNA cannot be ruled out as one of the mechanism involved in the AOAA-evoked neuronal loss.

Apart from *excitotoxicity*, AOAA displays potent convulsant properties. The compound evokes clonic seizures in rodents when applied systemically, intracerebroventricularly, or intrahippocampally (McMaster et al. 1991; Turski et al. 1991b, 1992). Diazepam, phenobarbital, and valproate, but not ethosuximide, trimethadione, acetazolamide, diphenylhydantoin, or carbamazepine, prevent convulsions induced by systemic administration of AOAA. Seizures evoked by intracerebroventricular injection of AOAA are blocked with KYNA and selective NMDA antagonists (Turski et al. 1991b, 1992).

Additionally, AOAA was demonstrated to induce seizures when injected into the rat entorhinal cortex (Du and Schwarcz 1992; Eid et al. 1995). In this paradigm, AOAA application resulted in selective neuronal loss within layer III of *entorhinal cortex* (Du and Schwarcz 1992). Notably, in contrast to striatum, the sequel of changes following application of quinolinic acid into the entorhinal cortex was qualitatively different from AOAA-evoked changes. Animals subjected to quinolinate did not display behavioral seizures; however, neuronal loss in layer V of the entorhinal cortex, often accompanied by neurodegeneration in the superficial layers of the dorsal perirhinal cortex, was a constant finding (Eid et al. 1995).

Novel data indicate that AOAA inhibits KYNA synthesis more potently in the milieu mimicking diabetic hyperglycemia. High level of glucose (30 mM) reduced the IC<sub>50</sub> value of AOAA by approx. 50 % (from 11.6 to 7.1  $\mu$ M), suggesting that severe hyperglycemia may render the brain more susceptible to the excitotoxic events (Chmiel-Perzyńska et al. 2007).

The above data suggest that AOAA is a valuable tool allowing detailed studies of excitotoxicity associated, at least in part, with deficient synthesis of brain KYNA.

## 6 KYNA and Brain Disorders

## 6.1 Parkinson's Disease

Parkinson's disease (PD) is one of the most common progressive neurodegenerative disorders, affecting approx. six million people worldwide. Although clinically heterogenous, PD is primarily characterized by tremor, bradykinesia, and muscle rigidity associated with dopaminergic denervation, primarily due to neuronal loss within substantia nigra pars compacta. Later on, other motor signs such as postural instability and impairment of gait and speech develop due to alterations in

non-dopaminergic transmission. Moreover, autonomic dysfunction and cognitive impairment arise in the course of disease (Lees et al. 2009). Familial forms of PD account for 10–15 % of diagnoses. Pathogenesis of remaining 85–90 % cases of PD is still unclear, and only some of the underlying mechanisms have been clarified so far. Disturbances of *ubiquitin-proteasome system*, dysfunction of mitochondria, and oxidative stress are among the recognized factors (Ali et al. 2011). Accumulating data indicate possible role of altered KYNA metabolism in the pathogenesis of PD (Szabó et al. 2011).

Consistently with neuroprotective as well as antiparkinsonian effect of glutamate receptors antagonists (Klockgether and Turski 1990; Klockgether et al. 1991; Turski et al. 1991a), application of KYNA or approaches aimed to increase its central level were shown to ameliorate neuronal loss and behavioral symptoms in experimental models of PD.

KYNA, injected intracerebrally, reduced akinesia, tremor, and rigidity in a severely parkinsonian monkeys subjected to *1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP)*, mitochondrial toxin commonly used to replicate PD-like symptoms in experimental animals (Graham et al. 1990). Parkinsonian symptoms are thought to result partially from the overactivation of neurons in the output areas of basal ganglia, i.e., substantia nigra pars reticulata and the medial segment of globus pallidus. In rat and primate models of PD, the direct injection of KYNA into the medial segment of the globus pallidus reduced behavioral changes (Brotchie et al. 1991). Similarly, in monkeys with hemiparkinsonism evoked by unilateral, intra-arterial MPTP application, the infusion of KYNA into contralateral globus pallidus internus partially alleviated symptoms of disease (Butler et al. 1997).

In rats, intraventricular administration of KMO inhibitor, *nicotinylalanine*, together with intraperitoneal kynurenine and probenecid infusion, evoked up to threefold increase of brain KYNA concentration (Miranda et al. 1997). Immunohistochemical assessment revealed that this rise of KYNA level was sufficient to protect nigrostriatal dopaminergic neurons from neurodegeneration caused by intranigral administration of quinolinic acid and to reverse some behavioral alterations (Miranda et al. 1997, 1999). Intraperitoneal treatment with kynurenine and probenecid prevented also neuronal damage and behavioral alterations induced by intrastriatal injections of *6-hydroxydopamine* (Silva-Adaya et al. 2011).

In primate model of MPTP-induced parkinsonism, orally administered *KMO inhibitor*, *Ro* 61-8048, increased central kynurenine and KYNA levels. This treatment was shown to improve L-DOPA-induced dyskinesias without alteration of drug antiparkinsonian efficacy (Samadi et al. 2005). Indeed, accumulating data suggest that glutamatergic manipulation of basal ganglia output is a promising approach to treat L-DOPA-induced dyskinesia in PD (Brotchie and Jenner 2011).

KYNA deficiency seems to play an important role in the development and progress of parkinsonian symptoms. An active metabolite of MPTP,  $MPP^+$  reduced cortical KYNA synthesis and inhibits KAT activity in vitro (Luchowski et al. 2002; Urbanska 2005). These results were confirmed in vivo, in a study demonstrating

that MPTP treatment decreases the number and optical density of KAT I immunoreactive neurons in substantia nigra of mice (Knyihár-Csillik et al. 2004). Similarly, application of 6-hydroxydopamine induced dose-dependent loss of KAT I immunoreactivity in the middle and lateral areas of substantia nigra pars compacta (Knyihár-Csillik et al. 2006). In human neuroblastoma cell line, pretreatment with KYNA attenuated MPP<sup>+</sup>-induced neuronal apoptotic cell death, decreased level of proapoptotic *Bax* protein, reduced MPP<sup>+</sup>-evoked mitochondrial dysfunction, and prevented activation of *caspase* 3 and 9 (Lee et al. 2008).

Human studies revealed decline of KYNA and kynurenine in frontal cortex, putamen, and substantia nigra of PD victims (Ogawa et al. 1992). Tryptophan/ kynurenine and kynurenine/KYNA ratios remained unchanged, and the therapy with L-DOPA did not seem to influence brain KYNA (Ogawa et al. 1992). In contrast, the level of KYNA in caudate and precentral cortical gyrus did not differ from control values, as shown by another study from the same group aimed to analyze brain KYNA content in Huntington's, Parkinson's, and Alzheimer's diseases (Beal et al. 1992).

In periphery, KYNA content within erythrocytes obtained from PD patients, as well as the activity of KAT II but not of KAT I, are increased (Hartai et al. 2005). Interestingly, the authors have also detected the activity of KAT I and KAT II in the serum of both patients and controls. In PD, the activity of both enzymes was decreased; however, there was no concomitant change in serum KYNA level (Hartai et al. 2005).

In the view of available data, deficient KYNA synthesis seems to be one of the underlying factors contributing to the pathogenesis of PD.

Surprisingly, there is no clinical data assessing potential efficacy of KYNA derivatives or NMDA antagonists in the treatment of dyskinesia in PD. Future research should be aimed to develop compounds acting selectively within basal ganglia. Targeting KYNA synthesis within specific brain areas seems to be one of the attractive therapeutic options.

### 6.2 Huntington's Disease

Huntington's disease (HD) is an inherited autosomal dominant neurodegenerative disorder associated with progressive loss of neurons, primarily in caudate-putamen. During later stages of disease, the cerebral cortex (particularly layers III, V, and VI), globus pallidus, thalamus, subthalamic nucleus, substantia nigra, white matter, and the cerebellum can also be markedly affected (Zuccato et al. 2010). Clinically, HD is manifested by progressive emotional, motor, and cognitive impairments. The development of HD results from *CAG trinucleotide* repeat expansion in the exon-1 of the gene encoding for huntingtin protein on chromosome 4. The length of CAG repeats correlates with age at onset of motor symptoms; however, HD patients may differ dramatically in disease manifestations, despite similar genetic changes. It is established now that multiple genes distinct from the HD locus itself may modify the course of disease (Zuccato et al 2010; Bano et al. 2011).

As with other human pathologies, proper animal models are invaluable tools that allow detailed studies of pathophysiological mechanisms underlying disease development. Initially, HD animal models were based on the intrastriatal injections of selective *excitotoxins*, such as kainate and quinolinic acid, indirect excitotoxin, AOAA, and peripheral application of mitochondrial toxin, 3-NPA (Coyle and Schwarcz 1976; Schwarcz et al. 1983; Urbanska et al. 1991; Brouillet et al. 1993). In rats, intrastriatal injection of AOAA leads to neuropathological and behavioral sequel mimicking HD (Urbanska et al. 1989, 1991; Beal et al. 1991) (see paragraph 5). Chronic, intraperitoneal administration of low doses of 3-NPA was demonstrated to induce relatively selective neuronal loss within striatum, accompanied by behavioral changes replicating symptoms of HD (Gould and Gustine 1982; Hamilton and Gould 1987; Brouillet et al. 1993).

It seems of interest that both AOAA and 3-NPA inhibit KATs activity, and, notably, AOAA is one of the most potent inhibitors of KATs (Luchowski et al. 2002; Urbanska 2005). 3-NPA was demonstrated to reduce the number of KAT I immunopositive glial cells in rat striatum and temporal cortex (Csillik et al. 2002). Susceptibility to 3-NPA action increases with age of animals (Csillik et al. 2002). Thus, results from chemical models of HD suggest that striatal deficiency of KYNA might be one of the factors contributing to the development of HD.

Following the discovery of genetic base of HD, novel animal models emerged. Disturbed function of kynurenine pathway was reported in heterozygous, symptomatic FVB/N mice transgenic for mutant *huntingtin* (Guidetti et al. 2000). Interestingly, transgenic animals showed large, over tenfold elevations in 3-hydroxykynurenine content in both striatum and cortex accompanied by smaller increases of KYNA levels, resulting in a prominent increase of the 3-hydroxyky-nurenine/KYNA ratio (Guidetti et al. 2000).

Human studies performed in later stages of disease revealed decline of CSF KYNA levels in HD (Beal et al. 1990; Heyes et al. 1994). In the brain tissue, twofold increase of KYNA in the motor cortex in Brodmann's area 4, but not within caudate, globus pallidus, and prefrontal cortex, was reported (Connick et al. 1988, 1989). Further studies demonstrated, however, that KYNA concentration in *putamen* and cortex of HD patients is decreased (Beal et al. 1990, 1992). Others have also reported approx. 50 % reduction of KYNA content in caudate, but not in hippocampus or cortical areas, in specimens obtained from late stage of HD (Jauch et al. 1995). In putamen, KYNA content reached approx. 67 % of control, however, at the border of statistical significance. Enzymatic analyses revealed decreased activity of KAT I and II (in putamen  $\sim$ 48–55 % of control) (Jauch et al. 1995). Notably, deficiency of KYNA ratio (Beal et al. 1990), indicating possible activation of the kynurenine pathway arm leading to neurotoxic metabolites.

Indeed, an increase of *3-hydroxykynurenine* in the brain was demonstrated by different groups (Reynolds and Pearson 1989; Pearson and Reynolds 1992; Guidetti et al. 2000, 2004). In specimens of putamen and temporal and frontal cortices, the concentration of 3-hydroxykynurenine was increased by nearly twofold

(Reynolds and Pearson 1989; Pearson and Reynolds 1992). Others reported that neostriatal and neocortical 3-hydroxykynurenine levels are increased by five- to tenfold, whereas KYNA was only slightly elevated in cortical samples obtained from early-stage HD patients (Guidetti et al. 2000). Further analysis of 3-hydroxykynurenine and quinolinic acid levels in the brains of patients with early or late HD has revealed an increase of these toxic kynurenine metabolites in 0/1 stage of disorder, but not during later phases (Guidetti et al. 2004).

Overall, an increased 3-hydroxykynurenine/KYNA ratio in HD patients and transgenic mice modeling the disease indicates the presence of abnormally high level of neurotoxic kynurenine metabolite accompanied by comparatively deficient formation of neuroprotective KYNA (Pearson and Reynolds 1992; Jauch et al. 1995; Guidetti et al. 2000, 2004). Based on accumulated data, it seems that changes in brain KYNA content and kynurenine pathway function are hallmarks of HD. In early stages of disease, both KYNA and 3-hydroxykynurenine levels may be increased. Later on, KYNA production seems to be severely compromised and 3-hydroxykynurenine/KYNA ratio is consistently higher (Pearson and Reynolds 1992; Jauch et al. 1995; Guidetti et al. 2000, 2004). Whether these changes reveal causative association or merely reflect consequences of progressive neuropathology remains to be established. However, accumulated data favor the opinion that disturbed kynurenine pathway function could be a component of pathological sequel contributing to the development of HD (Schwarcz et al. 2010).

Consequently, a number of studies were undertaken to evaluate the potential therapeutic benefits of increasing KYNA and/or decreasing 3-hydroxykynurenine level, via inhibition of KMO. The depletion of NADPH-diaphorase positive neurons caused by intrastriatal infusion of quinolinic acid was prevented by the intracerebral injections of nicotylalanine, an inhibitor of kynureninase and KMO, combined with intraperitoneal administration of kynurenine and probenecid, an inhibitor of organic acid transport (Harris et al. 1998). Chronic oral administration 2-(3,4-dimethoxybenzenesulfonylamino)-4-(3-nitrophenyl)-5-(piperidin-1-yl) of methylthiazole (JM6), small-molecule prodrug, acting as inhibitor of KMO, extended life span, prevented synaptic loss, and decreased microglial activation in the mouse model of HD (Zwilling et al. 2011). Similarly, in a fruit fly model of HD, KMO inhibition with UPF 648 [(15,2S)-2-(3,4-dichlorobenzoyl)-cyclopropane-1-carboxylic acid] provided neuroprotection against degeneration of photoreceptor neurons (Campesan et al. 2011). In transgenic mkat-2-/- mice, characterized by higher susceptibility to quinolinate-induced excitotoxic neuronal loss, application of UPF 648, KMO inhibitor, reduced quinolinic acid toxicity (Sapko et al. 2006).

Recently, a novel KYNA analogue, *N*-(2-*N*,*N*-dimethylaminoethyl)-4-oxo-1*H*quinoline-2-carboxamide hydrochloride, administered intraperitoneally, completely prevented the atrophy of the striatal neurons, increased survival time by about 30 %, and ameliorated hypolocomotion in transgenic *N171-82Q mice* (Zádori et al. 2011).

Thus, available data suggest that shift of kynurenine pathway toward KYNA synthesis may be neuroprotective in HD, whereas an excess of

3-hydroxykynurenine can be pathogenic. Moreover, changes in the levels of kynurenine pathway metabolites resulting in decrease of 3-hydroxykynurenine/ KYNA ratio seem to be of potential therapeutic value. It would be of great interest to evaluate the efficacy of KMO inhibitors, as well as modulators of other kynurenine pathway enzymes, in early-stage and advanced HD.

## 6.3 Alzheimer's Disease

Alzheimer's disease (AD) is the most common form of dementia characterized by the progressive memory loss and deterioration of higher cognitive functions. Neuropathological findings include the presence of extracellular senile (neuritic) plaques composed from various proteins, such as  $\beta$ -amyloid,  $\alpha$ -synuclein, or presenilins. Another hallmark of AD is the occurrence of intracellular neurofibrillary tangles containing hyperphosphorylated-tau protein and apolipoprotein E (Chintamaneni and Bhaskar 2012). The pathogenic process seems to start earlier than the onset of clinical symptoms, and it is estimated nowadays that neurodegeneration in AD begins decades before episodes of memory loss (Chintamaneni and Bhaskar 2012).

KYNA metabolism in the course of AD has been researched; however, the data are not conclusive. Initially, no change in the concentration of L-kynurenine and KYNA in the brains of AD victims was reported (Beal et al. 1992). In contrast, CSF KYNA was significantly lower in patients suffering from AD (Heyes et al. 1992). Others have demonstrated a significant rise of KYNA level in putamen (192%) and caudate nucleus (177%), but not in cortical specimens obtained from 11 patients with advanced form of AD (Baran et al. 1999). A trend for a decrease of kynurenine and 3-hydroxykynurenine in frontal cortex, caudate nucleus, putamen, hippocampus, and cerebellum of AD brain was also noted (Baran et al. 1999). In putamen and caudate, elevated KYNA correlated with an enhanced KAT I activity. A minor increase of KAT II activity was detected only in the caudate nucleus of AD subjects (Baran et al. 1999).

Peripheral KYNA status in AD was subject of two recent studies. Serum and erythrocytic KYNA were decreased by approx. 35 % in patients with AD, whereas kynurenine level and the activities of KAT I and KAT II remained unchanged (Hartai et al. 2007). The peripheral decline of KYNA accompanied by unaltered kynurenine levels was confirmed in study analyzing plasma samples of AD patients. Notably, quinolinic acid concentration was increased, suggesting an activation of kynurenine pathway arm leading to neurotoxic metabolites (Gulaj et al. 2010).

Only few reports analyzed the concentration of another neurotoxic kynurenine derivative, *3-hydroxykynurenine*, in the course of AD. Studies postmortem revealed an increase of 3-hydroxykynurenine content in temporal cortex of AD victims (Pearson and Reynolds 1992). In contrast, patients with AD exhibited decreased 3-hydroxykynurenine levels in CSF (Tohgi et al. 1992). Others found an increased

3-hydroxykynurenine/tryptophan ratio in the serum of AD patients, compared to age-matched and younger control (Widner et al. 2000).

In vitro studies have shown that fragments of amyloid peptide can stimulate IDO, thus activating an early step in kynurenine pathway what leads to an enhanced production of QUIN by human *macrophages* and *microglia* (Guillemin et al. 2003). Thus, existing data implicate the peripheral deficiency of KYNA in earlier stages of AD accompanied by shift of kynurenine pathway toward synthesis of neurotoxic 3-hydroxykynurenine. Considering the fact that most of the current data are based on the analysis performed during late stage of disease, when cerebral neurodegeneration is advanced, it seems that the central changes in kynurenine metabolism require further studies.

So far, potential usefulness of increasing brain KYNA as a possible therapy in early AD has been the subject of one study. Indeed, 4-month oral application of KMO inhibitor, *JM6*, blocked the development of spatial memory deficits in a transgenic mouse model of AD (Zwilling et al. 2011). JM6 also reduced anxiety-related behavior and prevented synaptophysin loss indicating its protective efficacy toward synapses; however, it did not alter the occurrence of A $\beta$  plaques (Zwilling et al. 2011).

On the other hand, reduction of NMDA-mediated transmission by excessive KYNA levels may contribute to the deterioration of cognitive functions and memory in patients with AD (Baran et al. 1999). In fact, plasma levels of KYNA correlate positively with cognitive functions assessment in Mini Mental State Examination and with the result of Hachinski Ischaemic Score (HIS) in AD patients (Gulaj et al. 2010).

In the view of above findings, disturbances of kynurenine pathway seem to be evident in AD; however, the nature of central KYNA changes and their role in AD-related structural and behavioral pathologies require further studies.

## 6.4 Ischemia

It is estimated that cerebral stroke affects approx. 15 million of people every year and is the second leading cause of mortality worldwide. Rapid occlusion of cerebral artery (ischemic stroke) or rupture of cerebral blood vessel (hemorrhagic stroke) evokes inadequate local supply of oxygen and glucose followed by an increase of extracellular glutamate levels, overactivation of glutamate receptors, and subsequent influx of excessive  $Ca^{2+}$  into neurons. Increased *intracellular*  $Ca^{2+}$  combined with oxidative stress finally lead to neuronal death (Mattson 2008).

Various experimental data indicated that blockade of NMDA receptors may reduce consequences of ischemia and injury of brain tissue; however, the therapeutic window seems to be very narrow and limited to several minutes post-insult (Ginsberg 2008; McIntosh et al. 1990; Muir and Lees 2003). Consequently, different manipulations increasing the brain level of KYNA have become an interesting alternative and were studied in various models of ischemia/anoxia.

In vitro, KYNA derivative, 7-Cl-KYNA, reduced neuronal damage in CA1 area of organotypic hippocampal slice cultures deprived of oxygen and glucose

(Newell et al. 1995). Other attempts demonstrated the efficacy of *KMO inhibitors* acting via reduction of 3-hydroxkynurenine and quinolinate synthesis, thus shifting kynurenine metabolism to the formation of KYNA. In organotypic hippocampal slice cultures, KMO inhibitors decreased hypoxia-/hypoglycemia-induced neuronal death in the CA1 region (Carpenedo et al. 2002).

In vivo experiments demonstrated that KYNA attenuates brain edema associated with anoxic-ischemic brain damage in neonatal rats (Simon et al. 1986). In 1-weekold rats subjected to hypoxia combined with ligation of the left carotid artery, peripheral administration of KYNA reduced brain damage (Andiné et al. 1988). Pretreatment with KYNA reduced the size of infarction and improved neurological outcome in the rat model of *focal ischemia* following middle cerebral artery occlusion (Germano et al. 1987). Peripheral administration of high doses of KYNA (1,000 mg/kg) markedly attenuated the hippocampal neurodegeneration in the gerbil model of global ischemia (Salvati et al. 1999). In the model of transient ischemia in gerbils, application of KYNA via microdialysis probe attenuated lactate accumulation and subsequent hippocampal damage (Katayama et al. 1992). KYNA, given systemically (500 mg kg<sup>-1</sup>, IP) or administered locally through the dialysis probe (10 mM), significantly reduced the forebrain ischemia-evoked release of glutamate and aspartate within rat striatum (Ghribi et al. 1994). In contrast, topical administration of KYNA onto the cortex in a four-vessel occlusion rat model did not reveal any changes in ischemia-evoked release of aspartate or glutamate (Phillis et al. 1999).

KYNA precursor, kynurenine, reduced infarct size in mice subjected to permanent focal cerebral ischemia produced by electrocoagulation of the distal middle cerebral artery and prevented the pyramidal cell loss in the CA1 area of the *hippocampus* in gerbils subjected to transient bilateral carotid artery occlusion (Gigler et al. 2007). Moreover, kynurenine markedly impaired the ischemia-evoked changes of spatial memory (Gigler et al. 2007). In the model of global brain ischemia, kynurenine, applied either pre- or early post-insult, reduced cortical neuronal loss (Robotka et al. 2008).

Neuroprotective properties of KYNA were also confirmed in an animal model of *heatstroke*. Treatment with KYNA (30–100 mg/kg, IV), 4 h prior to the heat stress, attenuated hypothalamic neuronal degeneration and apoptosis in rats. Authors suggest that systemic delivery of KYNA may attenuate not only neuronal degeneration but also multiorgan dysfunction in rats after heatstroke (Hsieh et al. 2011).

Increase of endogenous KYNA in the brain achieved through pharmacological blockade of KMO effectively protects against *excitotoxicity* (Miranda et al. 1997; Harris et al. 1998; Schwarcz and Pellicciari 2002). Administration of KMO inhibitors, causing 200–300 % rise of brain KYNA level, significantly reduced postischemic brain damage induced by bilateral carotid occlusion in gerbils or by middle cerebral artery occlusion in rats (Cozzi et al. 1999).

Alterations in endogenously produced KYNA were assessed in some models of ischemia. In cortical slices, KYNA synthesis was inhibited by hypoxia and glucose deprivation (Turski et al. 1989). Similarly, in *organotypic hippocampal slice cultures*, the lack of oxygen and glucose decreased the amount of KYNA

released into the incubation medium (Carpenedo et al. 2002). Only limited data come from in vivo animal models. In gerbils subjected to transient global ischemia, the endogenous levels of KYNA and the activity of its biosynthetic enzymes, kynurenine aminotransferases I (KAT I) and II (KAT II) in the hippocampus, were not altered 24 and 72 h after ischemic episode (Luchowska et al. 2003).

Interestingly, in patients at early (up to 24 h after infarct) stage of stroke, serum KYNA correlated positively with *D*,*L*-homocysteine which is considered an independent risk factor of atherosclerosis and an emerging risk factor in cognitive dysfunction and stroke (Urbanska et al. 2006). Following acute ischemic episode, an elevated kynurenine/tryptophan ratio suggested activation of kynurenine pathway (Darlington et al. 2007). Patients who died within first 3 weeks after stroke demonstrated increased serum KYNA level in comparison to subjects who survived (Darlington et al. 2007). Recent clinical study revealed that the kynurenine/tryptophan ratio, measured up to 7 days poststroke, correlated with infarct size and severity of clinical symptoms assessed with NIHSS (Brouns et al. 2010).

Disturbances of kynurenine pathway in the course of acute brain ischemia seem to have potentially important implications. It is not clear yet whether observed alterations reflect reactive, inflammation-linked activation of kynurenine pathway or develop prior to the inflammatory changes. Further studies, evaluating the ratio between neurotoxic kynurenine metabolite 3-hydroxykynurenine and neuroprotective KYNA during early and late phase of stroke, should clarify this issue. Nevertheless, encouraging experimental data suggest that increasing brain KYNA may be an interesting alternative in the therapy of early stroke.

## 6.5 Seizures and Epilepsy

Among kynurenine pathway metabolites, *quinolinic acid* is a well-described, neurotoxic agent able to evoke seizures in rodents, similarly to kynurenine itself (Lapin 1978, 1983). In contrast, KYNA displays clear anticonvulsant properties, as demonstrated by various experimental approaches in vitro and in vivo.

Pretreatment of brain slices with KYNA precursor, kynurenine, inhibited spontaneous epileptiform activity induced by the lack of  $Mg^{2+}$  (Scharfman and Ofer 1997). KYNA suppressed spontaneous epileptiform burst discharges in the *CA3* region of rat hippocampal slices induced by  $Mg^{2+}$ -free medium or picrotoxin (Stone 1988) and inhibited penicillin-induced synchronized afterdischarges in hippocampal slices from immature rats (Brady and Swann 1988). KYNA also depressed the amplitude and duration of  $Mg^{2+}$ -free bursts and reduced the frequency of penicillinevoked discharges (Schneiderman and MacDonald 1989).

In vivo, KYNA efficiently blocks convulsions evoked by quinolinate, produced along second arm of kynurenine pathway (Foster et al. 1984; Lapin et al. 1986). An increase of central KYNA levels, due to systemic administration of kynurenine, prevented the development of pentylenetetrazol-evoked seizures (Vécsei et al. 1992; Németh et al. 2004) and convulsions induced by the extract of poisonous

Atlantic coast mussels (Pinsky et al. 1989). Systemic or intracerebroventricular injection of KYNA inhibited seizures caused by the administration of bicuculline (Turski et al. 1990). Infusion of KYNA into the nucleus of the solitary tract reduced the severity of seizures induced by bicuculline or pentylenetetrazol (Walker et al. 1999). KYNA was also demonstrated to block the NMDA-evoked convulsions (Chiamulera et al. 1990). In WAG/Rij rats, genetic model of *absence epilepsy*, spontaneous spike-wave discharges were inhibited by KYNA administration. KYNA blocked also NMDA-induced potentiation of spike-wave discharges in this model (Peeters et al. 1994a, b).

Intracerebroventricularly applied KYNA displays protective effects in electrically kindled seizures (Thompson et al. 1988). Treatment with KYNA prior to the electrical *kindling* stimulus significantly reduced the rate of kindling in prepubescent and adult rats (Thompson et al. 1988). However, systemic administration of kynurenine or tryptophan did not protect rats against seizures induced by hyperbaric oxygen (Dale et al. 2000). KYNA had no effect on nicotine-induced seizures in mice (Kiś et al. 2000).

Interestingly, *antiepileptic drugs* may affect the synthesis of KYNA. Phenobarbital, felbamate, phenytoin, and lamotrigine increased KYNA production and activity of KAT I in vitro (Kocki et al. 2006). Moreover, high-fat and low-protein and low-carbohydrate *ketogenic diet*, leading to depletion of glucose and stimulation of ketone bodies synthesis, may also increase KYNA formation (Chmiel-Perzyńska et al. 2011). Exposure to profound ketosis (15 mM  $\beta$ -hydroxybutyrate) (BHB) with concomitant low glucose level (0.5 mM) increased synthesis of KYNA both in cortical slices and glial cultures. An increased activity of KAT I and II was detected in *glial cultures* but not in cortical slices (Chmiel-Perzyńska et al. 2011). Similar results were observed under in vivo conditions. In rats maintained on ketogenic diet, concentration of KYNA in hippocampus and striatum, but not in cortex, increased about two- to fourfold (Zarnowski et al. 2012).

Brain synthesis of KYNA is altered by a seizure episode, possibly due to the activation of glial cells (Wu et al. 2005). Reactive *astrogliosis* is characteristic for seizure models that are accompanied by neuronal loss in susceptible brain regions. Considering the fact that astrocytes are the main source of KYNA within the brain, increased KYNA level in the areas with prominent gliosis is not unexpected and may constitute defense mechanism.

Indeed, in the model of kainate-induced limbic seizures associated with widespread neuronal brain damage, the extracellular KYNA concentration in piriform cortex was not changed during the first 4 h of status epilepticus, but increased 1 month later, when astrogliosis was prominent (Wu et al. 1991). Various chemically induced seizures seem to increase KYNA level in dorsal hippocampus, starting in the first few hours after seizure development (Wu and Schwarcz 1996). Others have not confirmed hippocampal increase of KYNA during kainate-evoked seizures (Baran et al. 1995). However, 3 h after kainate injection, KYNA levels were raised in other brain areas, e.g., in piriform cortex, frontal cortex, amygdala, caudate, and putamen. The most prominent KYNA increases were observed in animals that developed status epilepticus (Baran et al. 1995). Contrasting data were obtained in models of kindled seizures, developing over long period and assumed to mimic *epileptogenesis* more adequately. In amygdalakindled seizures, KYNA level was reported to increase only in nucleus accumbens, but not in hippocampus, cerebral cortex, olfactory bulb, striatum, thalamus, tectum, cerebellum, or pons/medulla, as assessed 50 days after the last kindled seizure (Löscher et al. 1996). Moreover, the development of pentylenetetrazol-induced kindling was shown to be associated with the progressive decrease of KYNA in caudate-putamen, entorhinal cortex, piriform cortex, amygdala, and hippocampus (Maciejak et al. 2009). Changes in KYNA levels correlated with the stage of seizures (Maciejak et al. 2009). Similarly, deficiency of brain KYNA was revealed in a genetic model of absence seizures (WAG/Rij rats). In these rats, the level of KYNA in frontal cortex is lower, comparing to age-matched controls, at the time point when spontaneous seizures develop (Kamiński et al. 2003).

Human studies have also revealed alterations of central KYNA levels in epilepsy. Concentration of KYNA in CSF from epileptic patients was not changed, but serum KYNA in patients with intractable complex *partial seizures* was decreased (Heyes et al. 1994). Others have shown that in children with *West syndrome* or infantile spasm, the level of KYNA in CSF is lower, whereas among patients with grand mal seizures or occipital paroxysm, the CSF KYNA remains unaltered (Yamamoto et al. 1994, 1995). In the group of children with epileptic or febrile seizures, disturbances in circadian rhythm of kynurenine synthesis were found. The production of KYNA, kynurenine, 3-hydroxykynurenine, and xanthurenic acid was elevated during nocturnal hours in comparison to controls. The increase of KYNA and xanthurenic acid was not as high as of other kynurenines during epileptic seizures, whereas during febrile convulsions all kynurenines were increased by the same degree (Muñóz-Hoyos et al. 1997).

# 7 Conclusion

Systematic evaluation and interpretation of data indicates in general the beneficial effects of increasing central KYNA level in models of neuronal loss and seizures. Current state of research supports the notion that the selective blockade of excessive glutamate-mediated neurotransmission with exogenous compounds may be valuable in terms of neuroprotection; on the other hand, it might hamper processes of neuronal repair.

Promising experimental data indicate that novel, devoid of side-effects methods to manipulate glutamate transmission in humans should be sought. Endogenously produced KYNA is an attractive candidate for such studies. Sophisticated research adopting broader, multifactorial approach such as the development of drugs able to discriminate between brain structures or able to modify kynurenine metabolism selectively, in different cellular and subcellular compartments, is needed. Emerging therapies aimed to increase the level of neuroprotective KYNA or to reduce the brain expression of neurotoxic 3-hydroxykynurenine may become an important avenue in the treatment of neurodegeneration, ischemia, and seizures.

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# Neurotoxicity in Psychostimulant and Opiate Addiction

Ewa Niedzielska, Bartłomiej Rospond, Lucyna Pomierny-Chamioło, Anna Sadakierska-Chudy, and Małgorzata Filip

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#### Abstract

Psychostimulants and opioids evoke toxicity within the central nervous system (CNS), primarily linked to action of the substance (e.g., enhancement of synaptic concentrations of dopamine) or by indirect effects (increases in extrasynaptic glutamate levels, toxic metabolites). Associated with the development of drug addiction, the above mechanisms lead to the several pathological processes within the complex network of brain cell-to-cell

Laboratory of Drug Addiction Pharmacology, Institute of Pharmacology Polish Academy of Sciences, Kraków, Poland

e-mail: malgorzata.filip@uj.edu.pl; filip@if-pan.krakow.pl

E. Niedzielska • B. Rospond • L. Pomierny-Chamioło

Department of Toxicology, Faculty of Pharmacy, Jagiellonian University, Kraków, Poland

A. Sadakierska-Chudy Laboratory of Drug Addiction Pharmacology, Institute of Pharmacology Polish Academy of Sciences, Kraków, Poland

M. Filip (🖂) Department of Toxicology, Faculty of Pharmacy, Jagiellonian University, Kraków, Poland

interactions, such as alteration in neurotransmitter systems, dysregulation of energy metabolism, changes in structure of cellular membrane, apoptosis, generation of reactive species (RS), glial/microglial cell overactivation, excitotoxicity and prominent deficits in brain morphology (neuronal loss, axonal damage), alterations of transcription factors, immediate early genes and epigenetic mechanisms, as well as neurocognitive and neuropsychological impairments.

#### Keywords

Abuse • Addiction • Amphetamine • Animal studies • Cocaine • Dependence • Human studies • Neurotoxicity • Opiate

List of Abbreviations				
5-HIAA	5-hydroxyindoleacetic acid			
5-HT	Serotonin			
AMPA	α-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid			
BDNF	Brain-derived neurotrophic factor			
CART	Cocaine- and amphetamine-regulated transcript			
CAT	Catalase			
CNS	Central nervous system			
CpG	Cytosine-guanine dinucleotide			
CREB	cAMP response element binding			
CYP450	Cytochrome P450			
DNMT	DNA methyltransferase			
FADD	Fas-associated death domain			
FasL	Fas ligand			
fMRI	Functional magnetic resonance imaging			
GABA	γ-aminobutyric acid			
GPx	Glutathione peroxidase			
GSH	Glutathione			
HDAC	Histone deacetylase			
HMT	Histone methyltransferase			
HSP72	72-kDa heat shock protein			
HVA	Homovanillic acid			
MAO	Monoamine oxidase			
MDA	Malondialdehyde			
MDMA	3,4-methylenedioxymethamphetamine ecstasy			
miRNA	microRNA			
MRI	Magnetic resonance imaging			
MRS	Magnetic resonance spectroscopy			
NMDA	N-methyl-D-aspartic acid			
OPRD	Delta opioid receptor gene			
OPRK	kappa opioid receptor gene			
OPRM	mu opioid receptor gene			
PET	Positron emission tomography			
rCBF	Cerebral blood flow			

rCMR	Regional cerebral metabolic rate
ROS	Reactive oxygen species
RS	Reactive species
SNP	Single-nucleotide polymorphism
SOD	Superoxide dismutase
SPECT	Single photon emission computed tomography
TH	Tyrosine hydroxylase
TNF	Tumor necrosis factor
VMAT	Vesicular monoamine transporter

# 1 Introduction

Neurotoxicity can result from exposure to several exogenous factors and drugs of abuse are of such example. Psychostimulant and opiate intake is widespread, and their repeated exposure leads to either physical damage or disruption in the activity of the nervous system.

# 2 Psychostimulants and Opiates

Psychostimulants (e.g., cocaine and amphetamines, like methamphetamine, 3,4-methylenedioxymethamphetamine (MDMA=ecstasy)) are drugs that temporarily enhance the activity of the CNS and can induce the feeling of euphoria in humans, while in laboratory animals locomotor overstimulation, reinforcement/ reward, as well as subjective effects in a drug discrimination task are produced. The most prominent pharmacological mechanism of psychostimulants is the facilitation of dopaminergic neurotransmission; however, these drugs also enhance noradrenaline and/or serotonin (5-HT) activity (for a review, see Fleckenstein et al. 2007; Sulzer 2011). Psychostimulants can be characterized as "releasers" (amphetamines) and "uptake blockers" (cocaine) based on the mechanism of their acute effect on neurotransmitter flux through the monoamine transporter.

The direct actions of amphetamine and methamphetamine include the reversal of transport across the dopamine transporter (in lower extracellular concentrations; Sitte et al. 1998), diffusion into the cell due to its lipophilicity at higher extracellular concentrations (Sulzer et al. 1995; Kahlig et al. 2005), inhibition of the dopamine transporter, and inhibition of the vesicular monoamine transporter (VMAT)2 function; together, these actions induce stimulation-independent dopamine release (Sitte et al. 1998; Riddle et al. 2002). Amphetamine also enhances dopamine synthesis (Kuczenski 1975) and impairs dopamine metabolism by inhibiting monoamine oxidase (MAO) (Ramsay and Hunter 2002). In high concentrations or with repeated treatment, amphetamine inhibits tyrosine hydroxylase (TH) and/or TH protein levels (Ellison et al. 1978; Bowyer et al. 1998).

In contrast to amphetamine and methamphetamine, MDMA's mechanism depends on the animal species (Green et al. 2003). In nonhuman primates MDMA acts as a 5-HT neurotoxin; in rats, MDMA depletes 5-HT terminal markers due to enhanced 5-HT release (reviewed by Green et al. 2003); in mice MDMA releases dopamine (Rothman and Baumann 2003).

The primary mechanism of cocaine is linked to its ability to bind to the dopamine transporter, inhibit dopamine reuptake, and elevate synaptic and extrasynaptic dopamine levels (Brown et al. 2001), whereas it was more recently established that cocaine binding sites also contain both high- and low-affinity binding components (i.e., dopamine  $D_2$  receptors) (Marcellino et al. 2010), indicating a novel allosteric agonist action of cocaine in low (1–10 nM) concentrations on dopamine receptors (Ferraro et al. 2010). Apart from being a dopamine transporter blocker, cocaine also inhibits the noradrenaline and 5-HT transporters (Gatley et al. 1998). Of less significance is cocaine's action on the VMAT-2; cocaine enhances the storage of cytoplasmic catecholamines into synaptic vesicles, acting on the VMAT-2 which further allows for massive synaptic neurotransmitter release in response to depolarization (Brown et al. 2001).

It should be noted that metabolism of psychostimulants generates active substances that have their own toxic effects on nerves, including the brain. Thus, metabolites of amphetamine and methamphetamine include norephedrine, hydroxyamphetamine, hydroxynorephedrine N-hydroxymethylamphetamine, and possibly N-formylamphetamine (Jenkins and Cone 1998; Soloway and Patil 2011), by virtue of an action of cytochrome P450 (CYP450) on the parent drug. Metabolism of MDMA starts with O- or N-demethylation resulting in  $\alpha$ -methyldopamine, N-methyl- $\alpha$ -methyldopamine, or 6-hydroxy- $\alpha$ -methyldopamine, while during further biotransformation, conjugation with thiols or glutathione (GSH) leads to thioether derivatives or 5-(glutathion-S-yl)- $\alpha$ -methyldopamine; all the above metabolites are toxic to neurons (Milhazes et al. 2006; Capela et al. 2006; Cunha-Oliveira et al. 2008; Barbosa et al. 2012; Martinez et al. 2012).

Neurotoxic cocaine metabolites are benzoylecgonine, norcocaine, and cocaethylene (Nassogne et al. 1998; Bunney et al. 2001).

Opiates are any of the narcotics derived from the opium poppy plant. Together with other psychoactive drugs called "opioids," they share mechanisms of action, i.e., they bind to opioid receptors (see below). Opiates (e.g., morphine, heroin) and opioids (oxycodone, methadone, pentanyl, tramadol, pethidine, dextropropoxyphene, etc.) induce a feeling of well-being and a relief of pain, while their long-term repeated use leads to addiction and tolerance. This group of addictive drugs is characterized by a strong withdrawal syndrome that includes both somatic and affective components. In laboratory animals, opioids produce a biphasic locomotor response, consisting of an initial reduction followed by an increase in locomotor activity, discriminative stimulus, and reinforcement, whereas withdrawal syndrome includes a constellation of jumping, wet-dog shakes, teeth chattering, forepaw treading, forepaw tremors, and diarrhea. Opiates/opioids exert their actions by binding to three opioid receptors,  $\mu$ ,  $\delta$ , and  $\kappa$ , and the former two are involved in drug pleasurable effects due to the stimulation of dopaminergic neurons. Following opiates/opioids, enhanced dopamine extrasynaptic levels result from two points: at the level of the dopamine synapse and at the level of the afferents to the dopamine cell bodies (reviewed in Wise 1987; Sulzer 2011). The enhancement of dopamine release in the striatum may result from an indirect effect of opioids leading to the disinhibition of tegmental dopaminergic neurons by inhibiting  $\gamma$ -aminobutyric acid (GABA) interneurons or GABA terminals containing opioid receptors in the ventral tegmental area or by the direct activation of opioid receptors on accumbal cholinergic or GABAergic neurons (Sulzer 2011).

Opiates undergo several metabolic changes leading to assorted derivatives possessing toxic potential. Morphine is converted mainly by glucuronidation into morphine-3-glucoronide and morphine-6-glucoronide. However, oxidative demethylation and further oxidation via CYT450-dependent processes (principally via CYP3A4 and CYP2D6) may lead to N-hydroxymethylnormorphine, normorphine, and hydromorphone (Coller et al. 2009). Of note, morphine-3-glucoronide shows rewarding efficacy in mice (Vindenes et al. 2006).

Heroin (diacetylmorphine) undergoes rapid deacetylation to 6-monoacetylmorphine, which is approximately six times more potent than morphine, and further deacetylation to morphine (Sawynok 1986).

Pethidine is converted via CYT450 to a neurotoxic norpethidine, while dextropropoxyphene undergoes biotransformation into nordextropropoxyphene, a cardiotoxic agent.

A hallmark of psychostimulants and opioids (as well as other abused drugs), even having different mechanisms of action, is that they all impinge and increase dopamine neurotransmission within the mesolimbic circuitry of the brain (the so-called reward system); these effects are seen after acute (above) or chronic drug exposure (Filip et al. 2010; Sulzer 2011). The reward pathway is composed of dopaminergic neurons that project from the ventral tegmental area to several cortical and subcortical structures. The key structure that executes reward processes and is implicated in the addictive effects of the drugs is the nucleus accumbens, the terminus of the mesolimbic dopaminergic pathway. The nucleus accumbens is a regulator of motivation and learning processes (Di Chiara et al. 1999). It should be underlined that although impairment evoked by abuse of addictive drugs starts in the brain areas processing reward, the long-term drug intake disrupts the whole brain and leads to dysfunctions of emotions, motivations, learning, memory, executive control, and cognitive awareness. These functions are executed by the amygdala and hippocampus (memory and learning), the insula (conscious emotions), the dorsal striatum (habit forming learning), the orbitofrontal cortex (motivation/drive, salience attribution, motivation, and compulsive behaviors), the prefrontal cortex (executive control), and the cingulate gyrus (inhibitory control, attention, and impulsivity) (Volkow et al. 2003). Interestingly, in drug addiction there are recognized strong feedback mechanisms within the above brain structures with a special significance for the prefrontal cortex. In fact, in drug addicts, dysregulation of prefrontal cortex activity impairs its inhibitory coordination of other brain circuits, fixing the maintenance of drug habit. The above impairment behaviors are under control of glutamatergic neurotransmission (reviewed by Tzschentke and Schmidt 2003; Volkow et al. 2003; Kalivas 2004). Glutamategic neurotransmission is a cellular mediator of synaptic plasticity as well as learning and memory processes, and the current notion of addiction indicates that aberrant forms of drug-induced synaptic plasticity and learning in the brain drive compulsive and relapsing behaviors (Dacher and Nugent 2011).

# 3 In Vivo Evidences of Neurotoxicity

# 3.1 Human Studies

#### 3.1.1 Psychostimulants

# **Neurological and Psychiatric Impairments**

As reported in vivo, chronic use with amphetamines or cocaine has been implicated in cognitive deficits and psychological and anatomical impairments in human addicts. Among chronic drug users, methamphetamine abuse is associated with impairment of episodic memory, executive functions (e.g., novel problem solving), complex information processing speed, and decision-making abilities (impulsivity and risky decision-making) (Bechara and Martin 2004; Scott et al. 2007). Smaller impairments are present in motor skills (e.g., coordination) and language abilities (Caligiuri and Buitenhuys 2005; Scott et al. 2007). Other studies report a high level of psychiatric symptoms such as auditory/visual hallucinations, suspiciousness, and delusional thoughts in methamphetamine users (McKetin et al. 2006); such symptoms typically last only hours or diminish within a week of withdrawal, but prolonged symptom episodes have also been reported (McKetin et al. 2006). Abstinent methamphetamine abusers suffer also from mood disturbances like depression and anxiety (London et al. 2004) that may occur after the use episode, but may last for many months thereafter (Zweben et al. 2004).

Literature findings about influence of chronic MDMA use on neurocognition are equivocal. Some studies point out that only a "heavy" MDMA use can cause primary memory dysfunction (Gouzoulis-Mayfrank and Daumann 2006), while a low dose of MDMA (one or two tablets containing the usual amount of MDMA) has no relevant effects on human cognitive brain function (de Win et al. 2006; Jager et al. 2007). Moreover, there is a concern that memory deficits in MDMA users, although subtle, might be a risk factor for earlier onset and/or more severe decline of age-related memory deficits in later years, possibly due to hippocampal dysfunction (Gouzoulis-Mayfrank and Daumann 2009). Moreover, chronic MDMA users reported significantly increased levels of emotional instability, depressive mood, anxiety, aggression, impulsive features, and paranoid ideation (O'Leary et al. 2001; Gouzoulis-Mayfrank and Daumann 2009). It should be added that several factors like polydrug use, genetic factors (Gouzoulis-Mayfrank and Daumann 2009), or positive family history of psychiatric problems

(O'Leary et al. 2001) have been recognized within MDMA abusers, making a linkage between MDMA abuse and the drug-induced psychological problems to be established in further studies. Disturbances in working memory were reported also for cocaine-dependent subjects in which functional magnetic resonance imaging (fMRI) analyses during a working memory task showed impaired activation in the frontal, striatal, and thalamic brain regions (Cowan 2007; Moeller et al. 2010).

Chronic cocaine users suffer from diminished decision-making capacity, disrupted perception of motivational drive, and impairments in self-control (Goldstein et al. 2007a). Some researchers indicate that cocaine use by mothers during pregnancy increases the risk of significant cognitive (e.g., lower IQ, poorer language development, impaired attention), emotional (impulsivity), and motor deficits in the offspring (Arendt et al. 1999; Smith et al. 2001; Harvey 2004; Jones et al. 2004). Of note, some researchers do not report noticeable neurological or psychiatric abnormalities in children born of mothers who abused cocaine during pregnancy (Eyler et al. 2001; Rose-Jacobs et al. 2002).

#### Morphological Changes

Amphetamine users, apart from functional manifestation of drug-induced neurotoxicity (above), show prominent morphological brain changes. In fact, a loss of gray matter in the cingulate, paralimbic, and limbic cortices; noticeable smaller hippocampus volume; and hypertrophy of white matter were found in chronic methamphetamine users (Thompson et al. 2004). Additionally, MDMA users reveal reduction of multiple gray matter regions in the neocortical, bilateral cerebellum, and midline brain stem brain regions, potentially involved in some previously reported neuropsychiatric impairments (Cowan et al. 2003).

There are also morphological abnormalities in cocaine abusers seen as decreases in gray matter in the orbitofrontal cortex, age-related hyperintense areas of white matter regions in the cerebral and insular subcortex in fMRI scans (Franklin et al. 2002). Apart from the above brain regions, magnetic resonance imaging (MRI) tests performed on cocaine-dependent subjects (N = 40) showed also lower gray matter volumes in cerebellar hemispheres and thalamus (Sim et al. 2007). The authors suggest that cerebellar deficits may contribute to neuropsychological deficits and motor dysfunction frequently observed in cocaine-dependent subjects. Long-term consumption of cocaine may also influence white matter connections in the brain (Lane et al. 2010); deficits in white matter integrity have an inverse relationship with the length of abstinence from cocaine abuse in cocaine-dependent patients (Xu et al. 2010). Cocaine was also found to upregulate heat shock protein expression, leading to blood–brain barrier breakdown and brain edema formation (Sharma et al. 2009).

Recently reviewed by Derauf et al. 2009, infants, children, and young adults display some morphological abnormalities on MR-based imaging scans as cortical infarcts, pachygyria, schizencephaly, paraventricular hemorrhage, subependymal and periventricular cysts, as well as decreases in the volume of the corpus callosum, caudate–putamen or thalamus, and in cortical gray matter in both cocaine and methamphetamine abusers or their offspring.

# **Microglia Changes**

Chronic methamphetamine users showed an increased level of glial activation marker myoinositol in the frontal gray and left frontal white matter (Ernst et al. 2000; Chang et al. 2007; Sung et al. 2007; Krasnova and Cadet 2009). With use of positron emission tomography (PET), Sekine et al. (2008) indicated that methamphetamine-dependent individuals present a remarkable level of microgliosis in several brain regions (the midbrain, thalamus, striatum, insular, and orbitofrontal cortices); these changes were correlated inversely with the duration of the drug abstinence. Because the above observation contrasts with the findings in autopsied methamphetamine users' brains (see 4.1.1.2), further correlation analyses between the time needed for induction and progression of microglia activation are required.

#### Metabolic Changes

Human amphetamine addicts have serious brain metabolism impairments that are significantly altered in cortical and striatal areas (Volkow et al. 2001a; Wang et al. 2004). Recent papers reported reduced concentrations of N-acetylaspartate, being a marker of neuronal activity linked with attention control (Ernst et al. 2000; Sung et al. 2007; Licata and Renshaw 2010), and of creatine that serves as a metabolic integrity marker (Sekine et al. 2002) in the basal ganglia of methamphetamine abusers. Moreover, chronic methamphetamine users display raised striatal volumes and altered brain glucose metabolism in the orbitofrontal and limbic regions in the frontal gray and left frontal white matter (Ernst et al. 2000; Sung et al. 2007; Chang et al. 2007; Krasnova and Cadet 2009) as well as a higher glucose activity in the parietal cortex and lower activity in the thalamus and striatum (Volkow et al. 2001a).

Similarly to chronic methamphetamine, MDMA users showed decreased ratios of N-acetylaspartate to creatine which were associated with memory deficits (Reneman et al. 2001); however, other authors have not reported such differences (Daumann et al. 2004).

In vivo imaging findings on human cocaine users showed acute cocaine-evoked activation of the anterior cingulate (Daglish and Nutt 2003; Romero et al. 2010), while presentation of cocaine-associated cues resulted in the significant activation of metabolic activity of the anterior cingulate cortex region (Goldstein et al. 2007b). Local perfusion deficits were confirmed by PET and single photon emission computed tomography (SPECT) imaging in active cocaine abusers in the orbitofrontal cortex where the glucose metabolism was elevated and linked closely to changes in cerebral metabolism (Holman et al. 1991, 1993; Levin and Kaufman 1995; Levin et al. 1995; Volkow et al. 1991, 2004; Martinez et al. 2009), and this hypermetabolic response correlated with intense craving in cocaine abusers (Volkow and Fowler 2000) and persisted after months of detoxification (Volkow et al. 1988). PET studies confirmed MRI analyses about decreased metabolic activity in the orbitofrontal cortex and cingulate gyrus in the detoxified cocaine abusers (Volkow et al. 2004); such decreases were associated with reductions in striatal dopamine  $D_2$  receptors (see below). In another study in active cocaine

abusers in both genders, PET and 2-deoxy-2[<sup>18</sup>F]fluoro-D-glucose indicated a reduction in whole brain glucose metabolism in female and a trend to increase in male subjects during cocaine-cues video (Volkow et al. 2011). Children with prenatal cocaine exposure showed increased striatal creatine levels in magnetic resonance spectroscopy (MRS) study what suggested an impairment of energy metabolism in the brain of children exposed to this drug in utero (Fowler et al. 2007). The findings observed in stimulant abusers document brain metabolic changes in areas involved in reward circuitry and dopamine systems.

## **Changes in Neurotransmission**

PET and MRI imaging studies have revealed that amphetamines can lead to neurodegenerative changes in the brains of human addicts; these abnormalities include persistent decreases in the levels of dopamine transporters in the orbitofrontal cortex, dorsolateral prefrontal cortex, and caudate-putamen (McCann et al. 1998a; Volkow et al. 2001b, c; Sekine et al. 2003; McCann et al. 2008) and the lower density of 5-HT transporters in global brain regions (Sekine et al. 2006) in methamphetamine addicts. PET imaging identified also partial recovery of dopamine transporter binding in methamphetamine abusers during abstinence (Volkow et al. 2001c). The dopamine transporter reduction paralleled slower motor function and impaired episodic memory functioning, enhanced severity of psychiatric symptoms, and longer duration of methamphetamine use, while the reduction of 5-HT transporter may have a connection with increased aggressive behavior in methamphetamine abusers (Scott et al. 2007). It should be added that deleterious effects of methamphetamine such as N-acetylaspartate, creatine decrease, and microglial overactivity are enhanced in HIV-positive methamphetamine abusers (Chang et al. 2005), which indicate additive detrimental effect of coexisting HIV infection and methamphetamine abuse on human brain function.

Studies in MDMA users have reported enduring decrements in global brain 5-HT transporter binding (Ricaurte et al. 2000); however, neither a significant correlation between reductions in 5-HT transporter availability and extent of MDMA abuse nor improvements in markers for transporter during periods of drug abstinence have been demonstrated (Buchert et al. 2006; Thomasius et al. 2006). Of note, a separate SPECT imaging study in MDMA recreational users reported reduced striatal dopamine transporter binding (Reneman et al. 2002).

A growing number of evidence indicate enhanced glutamate transmission in many limbic brain areas in the development of amphetamine neurotoxicity (Abekawa et al. 1994; Stephans and Yamamoto 1994; Rocher and Gardier 2001; Cunha-Oliveira et al. 2008). Thus, reduction in glutamate and glutamine transporter systems has been described in the frontal gray matter of abstinent chronic methamphetamine abusers but not in active drug addicts (Ernst and Chang 2008) as an adaptation leading possibly due to reduced methamphetamine excitotoxicity (Ramonet et al. 2004). Since downregulation of glutamate and/or glutamine transporter levels is lowest at the beginning of abstinence ( $\leq 1$  month), they may serve as a factor involved in methamphetamine craving (cf. Ernst and Chang 2008).

Another amphetamine-derived psychostimulant, MDMA, evokes reduction in the metabolite of 5-HT, 5-hydroxyindoleacetic acid (5-HIAA), levels in cerebrospinal fluid in human abusers (Ricaurte et al. 1990; McCann et al. 1994, 1999; Bolla et al. 1998), and such change is regarded as a marker of the drug-induced toxicity. Confirmed by PET studies showing significant and dose-dependent depletion of 5-HT transporter levels in most cortical and subcortical brain regions of chronic MDMA users (McCann et al. 1998b; Semple et al. 1999; Reneman et al. 2001), the latter depletion was negatively correlated with the extent of abuse (McCann et al. 1998b; Ricaurte et al. 2000) and paralleled memory and learning impairment (Cowan et al. 2003).

PET studies have been also performed to reveal cocaine-induced abnormalities in brain dopamine neurotransmission, including long-lasting decreases in  $D_2$  receptor density in cocaine abusers (Volkow and Fowler 2000). These observations concerning  $D_2$  receptor level could be linked to reduced sensitivity of reward circuits to stimulation by natural rewards and increased risk for drug taking (Volkow et al. 2004). Moreover, detoxified cocaine abusers had still a marked decrease in dopamine release as measured by methylphenidate-induced decreases in striatal [<sup>11</sup>C] raclopride binding (Volkow et al. 1997), while the dopamine transporter density – elevated shortly after cocaine abstinence – was normalized with long-term detoxification (Malison et al. 1998). Interestingly, no difference in  $D_1$  receptor density was observed between cocaine-dependent subjects and matched controls (Martinez et al. 2009). Another PET scan report showed the low VMAT-2 availability in the striatum of cocaine abusers which the authors link to a compensatory downregulation of the dopamine storage vesicles in response to chronic drug abuse and/or a loss of dopamine terminals (Narendran et al. 2012).

## Manifestations of Genomic Mechanisms

The areas of the genome underlying amphetamine-type (cocaine and methamphetamine) substance use disorders are coming into light. In an American Indian community at high risk for substance dependence, analyses of multipoint variance component LOD scores failed to yield evidence of linkage for stimulant dependence for a panel of 791 microsatellite polymorphisms. However, the importance of "craving" as an important addiction phenotype was associated with regions on chromosomes 12, 15, and 18 in this user population. In more detail, the "craving" phenotype and "heavy stimulant use" were associated with a locus found on chromosomes 15q25.3-26.1 near the nicotinic receptor gene cluster while a loci for stimulant "craving" on chromosomes 12p13.33-13.32 and 18q22.3 (Ehlers et al. 2011). The single-nucleotide polymorphism (SNP) 118A>G in the mu opioid receptor gene (OPRM1) coding sequence was associated with methamphetamine psychosis, having latency less than 3 years from first drug intake in Japanese subjects (Ide et al. 2004). In cocaine addicts, an association with drug intake was shown with three promoter SNPs (-1793T>A, -1699insT, and -2044C>A) of OPRM1 gene (Hoehe et al. 2000), while a very recent large study identified association between the 17TT genotype and a cocaine addiction in HIV-positive and HIV-negative African addicts (Crystal et al. 2012). In a study of methamphetamine dependence/psychosis, an association was found with SNP rs2075572 (IVS2+691G>C) of the latter gene in Japanese subjects (Ide et al. 2004, 2006), but not with the SNP rs2234918 (921T>C) in exon 3 of the delta opioid receptor gene (OPRD1) (Kobayashi et al. 2006).

# 3.1.2 Opiates

# Neurological and Psychiatric Impairments

It is well-established that chronic exposure to heroin often results in cognitive deficits such as poorer performance in learning, cognition, and memory in human addicts (Cipolli and Galliani 1987; Guerra et al. 1987; Spain and Newsom 1991; Eisch et al. 2000). Other findings showed that methadone itself may contribute to cognitive impairment. In fact, greater impairment on verbal fluency, processing speed, response inhibition, visuospatial attention, cognitive flexibility, and set shifting was found in methadone-maintained patients than in abstinent groups. Also, in direct comparison of the methadone-maintained patients to former heroin users, similar impairment in executive function, visual memory, and impulsivity was found (cf. Martin et al. 2007). Methadone-maintained subjects rated the subjective value of a monetary reinforcement lower than the healthy comparison group, too (Martin-Soelch et al. 2001). Acquired brain injury (e.g., head trauma, nonfatal overdose) following long-term heroin use may contribute to variability in cognitive impairment. Interestingly, buprenorphine, another opioid used for maintenance treatment of former heroin abusers, was found to cause less psychomotor and cognitive impairment than methadone (Curran et al. 2001).

# Morphological Changes

In heroin addicts neuroimaging studies indicate morphological changes within the posterior limb of the internal capsule and sparing of subcortical U fibers (cf. Martin et al. 2007). Another study (Lyoo et al. 2006) reported gray matter reductions in prefrontal, insular, and temporal cortices in patients on methadone maintenance compared with age- and sex-matched healthy subjects.

# Metabolic Changes

Two studies have reported disturbances in frontal gray matter of patients maintained on methadone, buprenorphine, or heroin as the patients had lower concentration of N-acetylaspartate, as measured by MRS (Haselhorst et al. 2002; Yücel et al. 2007). MRI scans revealed diffuse white matter changes which paralleled widespread confluent vacuolar degeneration of the deep white matter, sparing of the brain stem and cerebellar white matter and extensive axonal injury in two patients with fatal drug-associated leukoencephalopathy (Ryan et al. 2005). Another in vivo functional assay indicates acute left-sided sensorimotor hemiplegia occurring acutely after a first inhaled dose of heroin by a 16-year-old male (Vella et al. 2003). The latter authors detected disturbances predominantly in parieto-occipital subcortical white matter and both ventral globus pallidus in MRS data and postulate defects in mitochondrial function as well as hypoxic injury in specific and limited areas of white matter due to acute heroin.

In SPECT scans conducted on opiate-dependent (methadone-maintained or heroin-using) subjects, a decrease in regional (the frontal and temporal cortex) cerebral blood flow (rCBF) was shown (Danos et al. 1998). A similar pattern of rCBF reduction has also been reported during the first week (Rose et al. 1996; Danos et al. 1998) or 10 weeks (Botelho et al. 2006) of detoxification, while Rose et al. (1996) have reported improved rCBF in the frontal cortex in patients rescanned at 3 weeks of withdrawal. Long-lasting neurobiological abnormalities in the regional cerebral metabolic rate (rCMR) in the anterior cingulate cortex were detected following a 6-month abstinence period in opiate addicts using fluorodeox-yglucose PET (FDG-PET) (Galynker et al. 2000).

## **Manifestation of Oxidative Stress**

In heroin addicts (N = 34) chronic opioid use evokes significant growth of oxidative stress (Kovatsi et al. 2010).

# Manifestations of Genomic and Epigenetic Mechanisms

There is a strong association between major genes of the opioid system and addiction to opiates. The most widely studied are mu, kappa, and delta receptors encoded by OPRM1, OPRK1, and OPRD1 genes (cf. Levran et al. 2012). Within two important SNPs (i.e., 17C>T (Ala6Val) and 118A>G (Asp40Asn)) out of the variants described in dbSNP in the OPRM1 gene coding sequence, only the SNP 118A>G has been linked with opioid addiction, and such observations come from Swedish (Bart et al. 2005), Chinese, European American (Drakenberg et al. 2006), and Indian patients (Deb et al. 2010). Other studies did not detect the above association (Franke et al. 2001; Glatt et al. 2007; Levran et al. 2009) nor association between the17T allele and opioid dependence in Indian (Kapur et al. 2007) and African (Crowley et al. 2003) addicts. Similarly, no association with opioid addiction has been reported in African and European American subjects analyzing SNPs -1793T>A, -1699insT, -1320A>G, 17C>T, 118A>G, as well as SNPs 540825 and 562859 (Crowley et al. 2003; Smith et al. 2005).

There are some evidences for association of OPRK1 SNPs 1051660 (Yuferov et al. 2004; Gerra et al. 2007) and of OPRK1 SNP rs6473797 (Levran et al. 2008) with opioid addiction. Recently, some papers indicated other polymorphism SNPs (e.g., rs2236861, rs2236857, rs3766951, rs204055, rs2236857, and rs2298896) for the OPRD1 gene in human addicts; associations with heroin addiction strongly depend on ethnic groups (cf. Levran et al. 2012).

Being a multifactorial and polygenic disorder, drug addiction does not conform to a simple Mendelian transmission pattern. Thus, epidemiological studies conducted on family, twin, and adoption suggest involvement in both genetic variance and environmental factors to increase susceptibility to addiction that neither act in isolation (Dick and Foroud 2003). Recent studies from the last 10 years demonstrated that posttranscriptional regulation of gene expression plays a key role in neurogenesis and synaptic plasticity (Puckett and Lubin 2011) and supports the role of epigenetic processes to control drug addiction (Robison and Nestler 2011; Wong et al. 2011). A separate study indicated also the occurrence of epigenetic mechanisms (i.e., an increase in OPRM1 DNA methylation) in lymphocytes of methadone-maintained former heroin addicts (Nielsen et al. 2009). Another recent study demonstrates that overall methylation in human blood DNA from opioid addicts was increased and cytosine-guanine dinucleotide (CpG)islands located in OPRM1 promoter were hypermethylated (Chorbov et al. 2011). Additionally, in sperm-derived DNA the methylation was increased at CpG sites, but global methylation was not affected (Chorbov et al. 2011). Hypermethylation of OPRM1 promoter can cause gene silencing, while methylation in sperm may suggest heritability of dependence phenotypes.

# 3.2 Animal Studies

#### 3.2.1 Psychostimulants

# **Behavioral Manifestation of Neurotoxicity**

Methamphetamine was shown to induce only subtle to moderate functional manifestation of neurotoxic-like alterations in behavior of laboratory animals (cf. Gouzoulis-Mayfrank and Daumann 2009). Thus, only up to 3 days after a neurotoxic methamphetamine administration (4 times  $\times$  10 mg/kg, every 2 h) is there a decrease in spontaneous locomotor activity in rodents (Timar et al. 2003). In more subtle motor tests, i.e., active avoidance performance and balance beam performance, residual deficiency was reported in rats after methamphetamine neurotoxic treatment (Walsh and Wagner 1992). The most affected by methamphetamine intake were learning and memory behaviors. Neurotoxic methamphetamine exposure, under regimens of  $4 \times 10$  mg/kg, every 2 h, or  $24 \times 1.67$  mg/kg once every 15 min, impaired novel object recognition and impaired learning in a multiple T water maze test, with no differences on spatial navigation or reference memory in the Morris water maze in rats (Herring et al. 2008). Disturbances in consolidation of learned place preference and impairments on a radial maze sequential learning task and on a novelty preference object recognition were reported after neurotoxic methamphetamine doses in rodents (Chat-Mendes et al. 2007; Gouzoulis-Mayfrank and Daumann 2009). Interestingly, in nonhuman primates, gradual rise in methamphetamine doses in an escalating dose regimen diminished social behavior on "injection" days, with aggression declining during the study; such effects disappeared after 3 weeks of drug abstinence (Melega et al. 2008).

More conflicting findings concern functional manifestation of MDMA neurotoxicity. Some studies reported enhanced anxiety or weak memory performance in rodents and monkeys exposed to MDMA (Winsauer et al. 2002; Melega et al. 2008) but normal or return-to-normal task execution after 2–3 weeks abstinence even with previous 33-week MDMA escalated doses (Melega et al. 2008). Other findings indicate no assessment of functional impairments like risk-taking behavior or anxiety after MDMA administration (Gouzoulis-Mayfrank and Daumann 2009); according to the latter studies, MDMA-treated animals do not present functional abnormalities despite persistent neurotoxicity reported postmortem.

Prolonged exposure to cocaine (sometimes even an acute drug injection) resulted in cognitive impairment and/or a decrease in social contacts in animals (Daza-Losada et al. 2009; Murnane and Howell 2011). Monkeys exposed to self-administered cocaine showed attentional impairments with a more specific cognitive control impairment, implicating orbitofrontal cortex dysfunction (Porter et al. 2011).

#### Changes in Neurotransmission

Several neuroimaging and microdialysis findings reveal abnormalities within brain neurotransmission due to chronic psychostimulant exposure. Thus, imaging studies (PET, SPECT, or fMRI) indicate neurotoxic effects regarding dopamine and 5-HT brain systems due to long-term psychostimulant exposure to animals (for a review, see Howell and Murnane 2011; Murnane and Howell 2011). Neurotoxic effects toward dopamine system alterations include decreases in dopamine synthesis (lower DOPA level) (Melega et al. 1997) and a drop in dopamine transporter densities (Villemagne et al. 1998; Melega et al. 2000; Hashimoto et al. 2007) in the striatum of monkeys after two i.m. injections with methamphetamine/amphetamine or following 3–4 i.m. injections with methamphetamine to nonhuman primate (Villemagne et al. 1998; Hashimoto et al. 2007).

Neuroimaging reports showed neurotoxic results of MDMA action on 5-HT brain system during noncontingent drug administration. In fact, an early PET imaging study in a baboon characterized the effects of MDMA on in vivo 5-HT transporter availability using MDMA (5 mg/kg s.c., once or twice a day for 4 days) that caused diminished 5-HT transporter density in the caudate–putamen and cortical regions (Scheffel et al. 1998; Szabo et al. 2002). Such decreases in 5-HT transporter availability, however, were not evidenced during contingent MDMA administration in rhesus monkeys (self-administration model) even during higher MDMA dosage (100–140 mg/kg/day) and repeated drug treatment (12 months) (Banks et al. 2008). Long-term (18 months) self-administered MDMA to rhesus monkeys evoked also no significant change in the VMAT availability following at least 2 months of drug abstinence, as assessed with PET neuroimaging, despite an evidence of attenuation of reinforcing properties of MDMA perhaps through neurotoxic effects of MDMA (Fantegrossi et al. 2004).

PET imaging studies in monkeys showed that chronic exposure to selfadministered cocaine resulted in a significant reduction in dopamine  $D_2$  receptor levels (Czoty et al. 2004; Nader et al. 2006) that persisted for up to 1 year of abstinence in some monkeys (Nader et al. 2006). These studies demonstrate that monkeys with long-term histories of cocaine use reliably display lower dopamine  $D_2$  receptor densities in ways that correlate with cocaine dose and duration of exposure (Moore et al. 1998; Nader et al. 2002).

Data from in vivo microdialysis in rodents indicated long-lasting reductions in dopamine and 5-HT brain levels following amphetamine derivatives, including

methamphetamine (Herring et al. 2008) and MDMA (Gudelsky and Yamamoto 2008). Apart from enhanced release of monoamines in the striatum, nucleus accumbens, and prefrontal cortex, MDMA also evoked the release of acetylcholine in the striatum, hippocampus, and prefrontal cortex and increased the extracellular concentration of energy substrates, e.g., glucose and lactate, in the rat brain. In contrast to the acute stimulatory actions of MDMA on the release of monoamines and acetylcholine, repeated MDMA administration in high doses resulted in a neurotoxicity to 5-HT axon terminals in the rat (for a review, see Gudelsky and Yamamoto 2008).

Repeated cocaine self-administration evokes significant alterations in extracellular monoamine and amino acid levels in rodents' brain. Many in vivo microdialysis studies (Pettit and Justice 1989; Pontieri et al. 1996; Lecca et al. 2007; Saddoris et al. 2011) including a very recent paper by Wydra et al. (2013) indicated that self-administered cocaine evoked extracellular accumbal dopamine and that its level was much higher in comparison to yoked rats given passive *i.v.* cocaine injections. Ferris et al. (2011), using a combination of cocaine self-administration and microdialysis in freely moving rats, showed a decrease in dopamine transporter sensitivity to cocaine in the nucleus accumbens core. Such a significant decrease in the ability of cocaine to inhibit the dopamine transporter was apparent after a single day of self-administration, lasting for several days following cessation of cocaine intake (Ferris et al. 2011); the latter finding confirms PET imaging studies showing that acute cocaine administration decreases the availability of the plasma membrane transporter for binding, even after cocaine is no longer blocking dopamine uptake as evidenced by a return of dopamine to its basal levels (Kimmel et al. 2012).

A few other reports demonstrated increases in extracellular GABA derived from neuronal sources after 3 weeks of withdrawal from repeated passive cocaine administration (Xi et al. 2003), but no changes in GABA levels during cocaine self-administration (Sizemore et al. 2000). Moreover, in animals extinguished from cocaine self-administration, a reduction in basal extracellular GABA levels was observed in the ventral pallidum, while the reinstatement of cocaine seeking was associated with an augmentation of such effects (Tang et al. 2005). Changes in dopamine and GABA neurotransmission during the maintenance phase mirror the motivational aspects of cocaine intake. Apart from dopamine and GABA, cocaine also regulates glutamate extracellular levels with small increases at a few isolated time points during the self-administration sessions (Sizemore et al. 2000; McFarland et al. 2003; Madayag et al. 2007; Miguéns et al. 2008) as well as decreases in basal extracellular levels of nucleus accumbens glutamate (Wydra et al. 2013). Such glutamatergic inputs are believed to be critical for the learning and maintenance of goal-directed responding, including drug-seeking behavior (cf. Koob and Swerdlow 1988).

#### Metabolic Changes

In vivo studies on animals confirm human PET imaging data (see above) that amphetamines evoke serious functional alterations in glucose metabolism. Thus, methamphetamine repeated treatment (1 mg/kg, every other day for 5 days) decreased 18-FDG uptake in the hippocampus and in the insular cortex in rats conditioned to receive methamphetamine in a distinct environment. Hypoactivation of glucose metabolism in the hippocampus, amygdala, and auditory cortex was also seen after acute MDMA (20 mg/kg or 40 mg/kg) to rats (Soto-Montenegro et al. 2007). On the other hand, another PET study indicated that rats showing strong drug-associated place preference displayed large increases in 18-FDG uptake in the dorsal and ventral striatum, cingulate cortex, and retrosplenial cortex which proves their involvement in the conditioned effects of amphetamines (Carrion et al. 2009).

Chronic use of cocaine is also associated with lasting alterations in brain metabolism. A recent study documents cocaine-induced changes in brain metabolic activity as a function of history of drug voluntary intake (Henry et al. 2010). The latter authors showed that acute i.v. cocaine treatment resulted in increases in glucose metabolism in the anterior cingulate and medial prefrontal cortex. Similar metabolic decrements were found by Lyons et al. (1996) who reported that an *i.v.* infusion of cocaine (1 mg/kg) markedly decreased glucose utilization in the orbitofrontal cortex, entorhinal cortex, nucleus accumbens, and parts of the hippocampal formation in monkeys; smaller decrements were observed in the caudate and anterior putamen, too. The progressive involvement of cortical and striatal areas as a function of extensive cocaine exposure has also been demonstrated in monkevs with  $[^{14}C]^2$ -deoxyglucose and PET method (Lyons et al. 1996; Porrino et al. 2004). Initial exposures to cocaine resulted in metabolic effects of cocaine contained primarily in the ventromedial regions of the prefrontal cortex, while following chronic exposure to cocaine self-administration, glucose metabolism expanded mainly within the striatum.

Interestingly, changes in brain metabolism lasted even after long-term drug abstinence, as evidenced in the study with 28-day cocaine withdrawal in rats trained to self-administer cocaine where significant increases in the cerebral blood volume in the medial prefrontal cortex, caudate–putamen, motor cortex, and the pedunculopontine tegmental nucleus were detected (Chen et al. 2011).

# Manifestation of Oxidative Stress

Supporting the role of oxidative stress in toxicity of amphetamines, several antioxidants (e.g., N-acetylcysteine, vitamin E) or overexpression of antioxidant enzymes (e.g., superoxide dismutase (SOD)) attenuates the neurotoxic action of amphetamines (Colado et al. 1997; Imam et al. 2001, 2000; Wan et al. 2006; Wu et al. 2007).

#### Manifestations of Epigenetic Mechanisms

Separate studies conducted on transgenic animals indicated that overexpression of histone deacetylase (HDAC)4 or HDAC5 decreases behavioral responses to cocaine (Kumar et al. 2005; Renthal et al. 2007), whereas genetic deletion of *Hdac5* enhanced mouse responses to the chronic effects of cocaine (Renthal et al. 2007).

# 3.2.2 Opiates

# **Behavioral Manifestation of Neurotoxicity**

As evidenced in many papers, chronic exposure to opioids can lead to disturbances in cognitive functions and evoke anxiety in rodents. Impairments in learning and memory processes were detected in long-term memory consolidation in mice when heroin was administered after the learning sessions (Castellano 1975, 1980; Florian and Roullet 2004). Similar results were shown in discriminative avoidance paradigms – radial-, Y-, and plus-mazes and in the Morris water maze after chronic administration of morphine (Spain and Newsom 1991; Pu et al. 2002) and also in the Morris water maze following a single methadone dose administered 15 min before training to rats (Hepner et al. 2002) or chronic methadone in mice (Tramullas et al. 2007). Such memory impairment effects have been linked with opioid-induced neurotoxicity through activation of apoptotic pathways (see below).

#### **Changes in Neurotransmission**

With neuromolecular imaging it was documented that rats given passive single (10 mg/kg) or repeated (10 mg/kg) doses of morphine had significantly increased extracellular dopamine and 5-HT levels in the nucleus accumbens; the increases above 90 % in monoamine levels were seen already after a second dose of morphine. Decreases in dopamine neurotransmission after acute or repeated dosing of morphine, and still significantly lower on the 5th day of morphine abstinence, were detected in SPECT scans in monkeys (Xiao et al. 2006). Another effective toxic target related to morphine abuse is glutamatergic neurotransmission. Thus, in rodents chronic morphine use modifies extracellular concentrations of glutamate in the nucleus accumbens (Sepúlveda et al. 2004), prefrontal cortex, and hippocampus (Guo et al. 2005). Recently, spectroscopic analysis in rats showed that morphine administration (10 mg/kg) twice a day for one, six, or ten consecutive days resulted in a reduction in glutamate, with no statistically proven changes in glutamine, myoinositol, taurine, or N-acetylaspartate in the prefrontal cortex (i.e., 10 days of consecutive morphine treatment), while glutamate and myoinositol levels were elevated in the hippocampus (Gao et al. 2007). Other in vivo findings demonstrate short-lasting adaptations in neuronal glutamate relief (Nicol et al. 1996), synaptic glutamate uptake (Xu et al. 2003), expression of glutamate transporters (Mao et al. 2002), and neuronal activities of glutamate dehydrogenase and GABA transaminase (Kaur and Kaur 2001) following consecutive morphine administration.

# 4 Postmortem Evidences of Neurotoxicity

# 4.1 Human Studies

# 4.1.1 Psychostimulants

#### Changes in Neurotransmission

In humans several evidences of pathological changes have been observed in brain monoaminergic systems following abuse of amphetamines. Thus, postmortem analyses in chronic methamphetamine users have shown a reduction in levels of dopamine nerve terminal markers in the caudate–putamen (Wilson et al. 1996; Moszczynska et al. 2004; Kitamura 2009) and reductions in 5-HT transporter levels in the orbitofrontal and occipital cortices (Kish et al. 2009). Moreover, VMAT-2 levels were remarkably reduced in some methamphetamine users, which may indicate a neurotoxic action of the drug resulting from disturbances in striatal dopamine terminal integrity (Kitamura 2009).

Regarding MDMA neurotoxicity in human addicts, there are some proof showing that MDMA led to the severe reduction of striatal 5-HT and 5-HIAA, assessed shortly (21-h) postmortem in the brain of an MDMA user who had taken MDMA regularly for 9 years (Kish et al. 2000), while another high-dose MDMA user had diminished 5-HT transporter level and TH activity in the brain (Kish et al. 2010).

For the development of amphetamine neurotoxicity to dopamine and 5-HT terminals, the occurrence of hyperthermia seems to be important. Physical activity, like dancing in the clubs or even staying in a warm environment which prevents proper heat dissipation, contributes significantly to the general morbidity and mortality in MDMA (Milroy et al. 1996; Armenian et al. 2012; Parrott 2012) or methamphetamine (Ishigami et al. 2003; Ago et al. 2006) users. Disruption of blood–brain barrier permeability and brain edema development in autopsy cases after acute fatal intoxication of methamphetamine (Ago et al. 2006; Inoue et al. 2006) or MDMA (Milroy et al. 1996) were found. Findings from forensic autopsy cases indicate that intake of a lethal dose of methamphetamine (N = 11) with the occurrence of hyperthermia immediately before death resulted in the 72-kDa heat shock protein (HSP72) expression in the microvessels and ependymal cells in the hippocampus (Kitamura 2009).

Long-term cocaine exposure in human addicts leads to neurotoxic effects on dopamine neurons seen as reductions in the total number of striatal dopamine  $D_2$  receptors (Little et al. 2009). Other necropsy findings on cocaine abusers revealed serious dysfunction of monoaminergic systems, seen as decreases in the levels of VMAT-2 (Little et al. 2003) or an elevation of  $\alpha$ -synuclein expression in midbrain dopamine neurons together with an increase of the 5-HT transporter in the striatum and the limbic system (for a review, see Büttner 2011). Elevation in  $\alpha$ -synuclein expression may indicate increases in synaptic vesicle trafficking or pools, or enhancement in neuronal plasticity, or even selective loss of dopamine neurons (reviewed in Trexler and Rhoades 2012).

#### Activation of Microglia

Some postmortem studies in active methamphetamine users point to a significant increase in the number of microglia throughout the caudate–putamen, although such an increase neither paralleled immunohistochemical evidence for reactive microgliosis (Kitamura 2009) nor proliferation of the marker of reactive microglia CR3.43 (human complement receptor 3) (Kitamura et al. 2010). Also, long-term cocaine exposure by human addicts evokes infiltration of phagocytic cells (activated macrophages and activated microglia) within the caudate–putamen (Little et al. 2009). Of note, cocaine-evoked activation of microglial cells releasing

proinflammatory cytokines in the white matter and subcortical regions seems to be related to damage of neuronal cells (cf. Büttner 2011).

## **Manifestation of Oxidative Stress**

The formation of reactive species (RS) and the resulting oxidative and/or nitrosative stress is a common toxicological pathway of amphetamines, and this plays an important role in the neurotoxicity observed in the brain after chronic exposure to the psychostimulants in human addicts (Mirecki et al. 2004; Fitzmaurice et al. 2006; Carvalho et al. 2012). Thus, a significant increase in levels of 4-hydroxynonenal and malondialdehyde (MDA), being products generated from lipid peroxidation, was found in the caudate nucleus and to a smaller extent in the frontal cortex in autopsied brains of chronic methamphetamine users (Fitzmaurice et al. 2006). Such increases in lipid peroxidation products were linked with a depletion in GSH concentration and with an enhancement in the levels of Cu, Zn-SOD activity in the caudate of methamphetamine users (Mirecki et al. 2004). These data suggest that methamphetamine exposure in humans leads to abovenormal formation of potentially toxic lipid peroxidation products in the brain, while increased Cu,Zn-SOD activity seems to reflect possible compensatory response to oxidative stress due to amphetamine administration.

#### Manifestations of Apoptosis

Up to now, there is no definitive proof that apoptosis accounts for psychostimulant neurotoxicity in human addicts. Thus, a necropsy analysis on the striatum of chronic METH users (N = 11) showed a lack of caspase-3 activation (Kitamura et al. 2007). Another very recent postmortem human brain report by Alvaro-Bartolome et al. (2011) on cocaine abusers (N = 10) demonstrated also no activation of extrinsic and intrinsic apoptotic pathways in the prefrontal cortex and lack of activation of caspase 3 while the Fas-associated death domain (FADD) receptor complex and cytochrome c were downregulated. These authors showed also enhanced degradation of nuclear PARP-1, a hallmark of apoptosis, which was postulated to result from the formation of reactive oxygen species (ROS) by cocaine metabolites due to the induction of contraregulatory adaptations or non-apoptotic (neuroplastic) actions by the psychostimulant.

#### Manifestations of Genomic and Epigenetic Mechanisms

An early report on brains from humans with a history of cocaine abuse showed marked reductions in the levels of enkephalin mRNA and mu opiate receptor binding within the caudate and putamen, while levels of dynorphin mRNA and kappa opiate receptor binding in the putamen and caudate were elevated (Hurd and Herkenham 1993). As suggested by the latter authors, an imbalance between euphoric (enkephalin and mu opiate receptors) and dysphoric (dynorphin and kappa opiate receptors) markers may be linked with craving phenomenon detected in the human cocaine addict brain.

More recent microarray analysis showed alterations in brain gene expression in the postmortem anterior prefrontal cortex from human cocaine abusers (Lehrmann et al. 2006). Thus, the drug-specific changes included reduction in calmodulin-related transcripts (CALM1, CALM2, CAMK2B), while transcripts related to cholesterol biosynthesis and trafficking (FDFT1, apolipoprotein L = APOL2, SCARB1) and Golgi/endoplasmic reticulum functions (semaphorin 3B = SEMA3B, GCC1) were increased. As these changes represent common molecular features of drug abuse, they may underlie changes in synaptic function and plasticity that could have important ramifications for decision-making capabilities in drug abusers (cf. Lehrmann et al. 2006). Another gene array analysis in brains from cocaine abusers detected altered expression of some myelin-related genes in the nucleus accumbens, while in situ hybridization to quantify transcript expression of these genes showed only decreased expression of the transcript encoding PLP1. PLP1 is expressed at very high levels in oligodendrocytes and is essential in maintaining stability of myelin sheets, thus implicating adaptive changes following chronic cocaine abuse. Of note, postmortem expression of claudin-11 and transferrin was decreased in the caudate and internal capsule, respectively, in cocaine addicts (Kristiansen et al. 2009).

There are some early evidences about the epigenetic mechanisms due to psychostimulant intake. Chronic cocaine abuse induces alterations in gene expression in the nucleus accumbens of humans (Albertson et al. 2004). Microarray analysis revealed further that expression of cocaine- and amphetamine-regulated transcript (CART) was increased, but some myelin-related genes were decreased. Another study on human postmortem brain revealed transcription and epigenetic changes in the hippocampus (Zhou et al. 2011) with suppression of more than one third of genes responsible for ATP synthesis and increases of the transcript level of DNA methyltransferase (DNMT)3a, cAMP response element binding (CREB)1, HDAC2, and HDAC4. In addition, changes in H3K4me3 distribution were noted near promoters of active genes and away from promoters of protein-coding genes.

# 4.1.2 Opiates

#### **Changes in Neurotransmission**

Chronic opiate exposure impairs brain monoaminergic neurotransmission in human addicts. As evidenced in postmortem nucleus accumbens of 9 chronic heroin users, levels of TH protein and dopamine were reduced. On the other hand, striatal levels of the VMAT and 5-HT were normal (Kish et al. 2001).

#### **Changes in Protein Levels**

A recent postmortem report indicated that in the hippocampi of 29 lethally intoxicated heroin addicts, there was a significant increase in polysialic acid neural cell adhesion molecule, being one of the key players involved in plasticity and regeneration, of neural-tissue-positive neurons and glial cells, possibly reflective of an attempt to repair cell damage due to long-term heroin neurotoxicity and impaired plasticity (Weber et al. 2006). This may account for cognitive deficits seen in human addicts (see above).

In the brain of young opiate abusers (N = 34, all <40 years age), an excess of AT 8-positive neurofibrillary tangles was found in the subiculum, temporal neocortex,

nucleus basalis of Meynert, and the locus coeruleus, while beta-amyloid precursor protein is increased in the brain stem and temporal lobe (Ramage et al. 2005). The above changes may be risk factors for poorer performance in cognition and memory in opiate addicts; however, the direct proof is still lacking.

Another study performed on brains of chronic opiate addicts (N = 17) who had died of a heroin or methadone overdose showed a marked (47 %) reduction in levels of immunoreactive NF-L proteins in the frontal cortex (García-Sevilla et al. 1997); this change was considered as a specific damage of axonal transport resulting from long-term opiate-evoked neuronal damage.

#### Manifestation of Oxidative Stress

A study performed on samples of 17 human brains obtained 2–4 days after heroin users' death indicates high protein oxidation levels in all lobes of the cortex, brain stem, hippocampus, and white matter and lower peroxyl radical-trapping capacity – a marker reflective of the relationship between ROS amount and antioxidant efficiency (Gutowicz et al. 2006). The latter authors found an elevation of MDA levels, a lipid peroxidation product, in all parts of the brain except the hippocampal formation. Despite the above evidence for neurotoxicity, there was no reduction in the activity of antioxidant enzymes such as SOD or catalase (CAT) (Gutowicz et al. 2006).

#### Manifestations of Epigenetic Mechanisms

A significant change in the expression of numerous genes encoding proteins involved in presynaptic release of neurotransmitter was reported by Albertson et al. (2006) in heroin abusers. Thus, the transcript encoding the endogenous opioid peptide prodynorphin (*PDYN*) was decreased, while the neurotrophin receptor *NTRK2* (also known as *TrkB*) was significantly upregulated in heroin abusers. In addition, alterations in many genes involved in processes related to neurotransmitter release and recycling, including SNARE proteins, vesicular proteins, and recycling proteins, imply synaptic dysfunction in the brains of human heroin abusers. A very recent report by Michelhaugh et al. (2011) identified alterations of long noncoding RNA (lncRNA)s (such as MIAT, MEG3, NEAT1, and NEAT2) in heroin abusers compared to control subjects.

# 4.2 Animal Studies

#### 4.2.1 Psychostimulants

#### Changes in Neurotransmission

Amphetamines produce long-term decreases in markers of biogenic amine neurotransmission (mainly dopamine and 5-HT), and these neurotoxic effects include the decrease in monoamine levels and their major metabolites, reduced level of monoamine transporter binding sites, diminished expression and/or activity of enzymes involved in the metabolism, and synthesis of brain monoamine neurotransmitters – evident in rodents and nonhuman primates (Yamamoto et al. 2010; Carvalho et al. 2012). Thus, dopamine brain system impairments relate to decreased level of the neurotransmitter (Wagner et al. 1980; Ouinton and Yamamoto 2006), the dopamine metabolite, homovanillic acid HVA (Preston et al. 1985), reduced TH activity (Ellison et al. 1978; Hotchkiss and Gibb 1980), and a lower density of the dopamine transporter (Wagner et al. 1980; Krasnova et al. 2001) following high (usually 5–15 mg/kg in rats) doses and/ or longitude methamphetamine/amphetamine administration in rodents (Morgan and Gibb 1980; Ricaurte et al. 1982; Gouzoulis-Mayfrank and Daumann 2009) or nonhuman primates (Preston et al. 1985; Woolverton et al. 1989). In a model of a prevalent human methamphetamine exposure, produced by daily multiple doses in monkeys (escalation dosing of methamphetamine over 33 weeks, with final dosages resulting in estimated peak plasma methamphetamine concentrations of  $1-3 \mu$ M), postmortem analysis of brains showed persistent reductions in striatal dopamine levels and dopamine transporter binding associated with neurotoxicity in presynaptic dopamine system integrity (Melega et al. 2008). Similar long-term reductions in brain dopamine axonal markers were observed in baboons treated with methamphetamine (0.5-2 mg/kg, i.m., 4 times at 2 h intervals) (Villemagne et al. 1998).

On the other hand, the long-term (1.5 years) exposure to self-administered cocaine in monkeys evokes significant increases in dopamine transporter binding sites at the caudal level of the subcortical areas (the ventral caudate, ventral putamen, and nucleus accumbens) (Letchworth et al. 2001). The latter finding – showing similarity to studies in human cocaine addicts – suggests importance of this biomarker to the neurobiological effects of cocaine. Other studies on monkey i.v. self-administered cocaine for 18–22 months showed lowered dopamine  $D_2$  binding site density in the striatum (Moore et al. 1998); these data parallel studies on rodents and humans (see above).

The neurotoxic effects of amphetamines also apply to 5-HT-ergic nerves in the brain. Methamphetamine, although more toxic to DA neurons, leads also to 5-HT system impairments including (1) depletion of 5-HT and its metabolite 5-HIAA levels, (2) diminished tryptophan hydroxylase activity in the striatum and hippocampus, and (3) lower 5-HT transporter densities in rat frontal cortex, amygdala, and striatum (Hotchkiss and Gibb 1980; Ricaurte et al. 1980; Gouzoulis-Mayfrank and Daumann 2009). Another amphetamine derivate, MDMA, causes mainly 5-HT system impairments in rats or nonhuman primates (Quinton and Yamamoto 2006; Gouzoulis-Mayfrank and Daumann 2009; Carvalho et al. 2012) with reduction of 5-HT and 5-HIAA levels, decreased activity of tryptophan hydroxylase, and a diminished density of 5-HT transporter in rat cortical and subcortical brain areas (Battaglia et al. 1987; Itzhak and Achat-Mendes 2004; Cadet et al. 2007; Carvalho et al. 2012). Nevertheless, in mice, MDMA causes primarily dopaminergic neurotoxicity (Green et al. 2003), which demonstrates that not every species has the same sensitivity to amphetamines' actions. According to many studies, nonhuman primates seem to be more sensitive to the neurotoxic effects of MDMA than rodents, resulting in higher rates of 5-HT depletion with smaller doses of MDMA, and persisting hypoinnervation patterns in most neocortical

regions and the hippocampus examined even 7 years posttreatment (Ricaurte et al. 1992; Gouzoulis-Mayfrank and Daumann 2009). In fact, 18 months of MDMA selfadministration in primates produced 40–50 % depletion in 5-HT levels in the frontal, parietal, and temporal cortex, while lesser evoked downward trends in 5-HT depletions (approximately 25 %) were apparent in the occipital cortex, thalamus, and hypothalamus (Fantegrossi et al. 2004).

Methamphetamine was found to increase extracellular glutamate level followed by *N*-methyl-D-aspartic acid (NMDA) receptor activation and subsequent increase in intracellular Ca<sup>2+</sup> concentration (Cunha-Oliveira et al. 2008). As described earlier, a rise in the Ca<sup>2+</sup> concentration leads to the massive production of nitric oxide that in reaction with O<sub>2</sub> forms ONOO<sup>-</sup> (Capela et al. 2009); both NO and ONOO<sup>-</sup> lead to nitration of proteins including biosynthetic enzymes of dopamine and 5-HT (Kuhn and Arthur 1997; Kuhn et al. 1999; Carvalho et al. 2012). As hypothesized by Yamamoto et al. (2010), the oxidation and nitration of proteins associated with dopamine and 5-HT terminals appear to be the main mechanisms by which the amphetamine-evoked neurotoxicity arises.

Several studies have addressed the influence of hyperthermia following administration of amphetamines on neurotoxicity in animals. As found, methamphetamine-induced hyperthermia may be a factor increasing the drug-induced neurotoxicity, but it is not a necessary and sufficient condition in neurotoxicity induced by methamphetamine (Kita et al. 2003). Furthermore, hyperthermia might interact with other known mediators of methamphetamine neurotoxicity, such as increased glutamate neurotransmission, to enhance toxicity (see above).

#### **Metabolic Changes**

Some changes have been found in brain energy metabolism in rodents after amphetamine treatment. For example, subchronic amphetamine treatment (12-15 mg/kg/day for 7 days followed by 7-daily injections with 6-7 mg/kg/day)in rats caused enhanced local cerebral glucose utilization in the nucleus accumbens (Orzi et al. 1983). Increases in glucose metabolism were reported also after acute amphetamine administration (1-5 mg/kg) in extrapyramidal motor systems in rats (Wechsler et al. 1979; Porrino et al. 1984). Conversely, methamphetamine administration (4  $\times$  12.5 mg/kg, every 2 h) decreased local cerebral glucose utilization in the extrapyramidal system, also in the hippocampus and dorsal raphe nucleus in rats (Huang et al. 1999). These findings clearly demonstrate that amphetamines lead to regional, drug-dependent, and regimen-dependent alterations in brain metabolism.

#### Morphological Changes

There are several proofs that chronic treatment with psychostimulant drugs of abuse results in neuronal degeneration. Thus, in rats treated continuously with amphetamine (e.g., pellets implanted *s.c.* releasing amphetamine for 10 days) or methamphetamine (e.g.,  $10 \text{ mg/kg} \times 4$  times with 2 h intervals or 15 mg/kg s.c. every 6 h for 24 h), anatomical alterations in the form of swollen, dystrophic, or fragmented dopamine axons in the striatum have been found (Ellison et al. 1978; Hanspeter 1981; Ricaurte et al. 1982; Broening et al. 1997). Degenerations of presynaptic

5-HT axon terminals induced by high and/or repeated doses of methamphetamine (Axt and Molliver 1991; Cadet et al. 2003; McCann and Ricaurte 2004) were found throughout the forebrain of rats (O'Hearn et al. 1988), while in squirrel monkeys a reduction in 5-HT innervation was seen 7 years after exposure to the drug (Hatzidimitriou et al. 1999).

### **Glia and Microglia Activation**

As a consequence of amphetamine intake, damage of striatal axonal terminals was accompanied by reactive astrocytosis (Bowyer et al. 1994) and microglial activation (Pubill et al. 2002) via an increase in the expression of proinflammatory molecules (e.g., proinflammatory cytokines, interleukins). Interestingly, even an acute exposure to methamphetamine increased mRNA levels of IL-6, IL-1 $\alpha$ , and tumor necrosis factor (TNF)- $\alpha$  in mouse striatum (Sriram et al. 2006; Gonçalves et al. 2008). Neurotoxic methamphetamine exposure (4 times × 10 mg/kg, every 2 h, or 24 times × 1.67 mg/kg once every 15 min) resulted in an increase in glial fibrillary acidic protein level 3 days posttreatment (Herring et al. 2008). Methamphetamine alone or its metabolites might activate glial cells by virtue of elevated cellular dopamine auto-oxidation, resulting in profound formation of dopamine o-quinones, strong microglial activators (Kuhn et al. 2006; Yamamoto et al. 2010; Carvalho et al. 2012).

The ability of MDMA to induce microglial activation is more ambiguous. Thus, microgliosis induced by MDMA administration was found in the male (but not female) mouse striatum and was absent in the rat cortex or striatum. MDMA injections increased also production of the proinflammatory cytokine IL-1 $\beta$  in rat frontal cortex (Orio et al. 2004), while the *i.c.v.* administration of IL-1 $\beta$  intensified MDMA-induced 5-HT toxicity in the cortex. On the other hand, MDMA has been shown to have an immunosuppressive power that involves an increase in IL-10 production, an anti-inflammatory cytokine (Yamamoto et al. 2010).

#### Manifestations of Oxidative Stress

There is reliable evidence of oxidative damage in animal brain after repeated exposure to amphetamines (see Table 1). Thus, amphetamine, methamphetamine, or MDMA do increase lipid peroxidation (Jayanthi et al. 1998; Gluck et al. 2001; Alves et al. 2007; Miranda et al. 2007) and induce various changes in the levels of GSH (Harold et al. 2000; Carvalho et al. 2001; Frenzilli et al. 2007; Miranda et al. 2007) or activity of several antioxidant enzymes (Jayanthi et al. 1998; Frey et al. 2006; Frenzilli et al. 2007; Miranda et al. 2007), providing evidences for the importance of RS in amphetamine-induced toxicity.

Interestingly, the bidirectional changes in brain SOD activity were demonstrated after amphetamine or its derivatives. For example, subchronic amphetamine administration caused a rise in the SOD activity in rat hippocampus (Frey et al. 2006), while MDMA led to the drop in SOD activity in mouse hippocampus (Frenzilli et al. 2007). The stronger oxidative potential following MDMA derives from its highly reactive quinone intermediate metabolites (cf. Capela et al. 2007). Production of RS appears to be linked in large part with amphetamine-induced thermal instability

D-Amph, chronic passive administration		Dose and treatment regimen	Biomarker	Change (brain structure)	Reference
administration	Rat	1 mg/kg, 2 mg/kg or 4 mg/kg (i.p.)	SOD	$\uparrow$ (HIP), $\downarrow$ (STR) for 1 mg/kg	Frey et al. (2006)
		for 7 days		↑(HIP), ↓(STR) for 4 mg/kg	1
			CAT	$\uparrow$ (PFCx), $\uparrow$ (HIP) for 4 mg/kg	
				↓(HIP), ↓(STR) for 2 mg/kg	
D-Amph, chronic passive	Rat	20  mg/kg (s.c.) for 14 days	GST	↑(HL)	Carvalho et al.
administration			GPx	↑(STR, NAc, mPFCx)	(2001)
			GR	↑(HYP, PFCx)	
			CAT	$\uparrow$ (PFCx)	
METH, acute passive	Mouse	1 mg/kg ( <i>i.p.</i> ), $4 \times$ every 2 h	SOD	↓(FCx)	Jayanthi et al. (1998)
administration			CAT	↓(STR)	
			GPx	↓(FCx, STR)	
			MDA	↑(FCx, STR)	
METH, binge passive administration	Rat	10 mg/kg ( <i>i.p.</i> ), $4 \times \text{every } 2$ h	MDA	↑(STR)	Yamamoto and Zhu (1998)
METH, binge passive	Rat	20 mg/kg ( <i>i.p.</i> ),	GSH	↓(STR)	Moszczynska et al.
administration		$4 \times \text{every } 2 \text{ h}$	GPx	Ø (STR)	(1998)
			GR	Ø (STR)	1
			GGTP	Ø (STR)	
			G6PDH	Ø (STR)	
METH, binge passive	Rat	10 mg/kg ( <i>i.p.</i> ),	GSH	↑(STR)	Harold et al. (2000)
administration		$4 \times \text{every } 2 \text{ h}$	GSSG	↑(STR)	
METH, binge passive administration	Mouse	10 mg/kg (i.p.), $4 \times$ every 2 h	3-NT	↑(STR)	Imam et al. (2000)
METH, binge passive administration	Mouse	10 mg/kg ( $i.p.$ ), 4 × every 2 h	PC	↑(STR, HIP)	Gluck et al. (2001)

Table 1 (continued)					
Model treatment	Animal	Dose and treatment regimen	Biomarker	Biomarker Change (brain structure)	Reference
MDMA, binge passive	Adolescent	Adolescent 10 mg/kg $(i.p.)$ ,	MDA	1) (Mitochondrial fraction	Alves et al. (2007)
administration	rat	$4 \times \text{every } 2 \text{ h}$	PC	$\uparrow \int of$ whole brain)	
MDMA, subchronic passive	Mouse	2.5  mg/kg  (i.p.),  for  5  days	GSH	(HIP)	Frenzilli et al. (2007)
administration			SOD	(HIP)	1
MDMA, subchronic passive	Rat	5 mg/kg ( <i>i.p.</i> ),	GPx	(HIP)	Miranda et al. (2007)
administration		for 7 days	GSH	Ø (HIP)	1
			MDA	↑ (HIP)	1
Abbreviations: $\uparrow$ increase, $\downarrow$ dec	rease, $\emptyset$ no effe	Abbreviations: f increase, d decrease, d no effect, p-Amph p-amphetamine, CAT catalase, FCx rontal cortex, G6PDH glucose-6-phosphate dehydrogenase, GGTPelutamylranessentidase GSH elutathione GSCH reduced elutathione GST elutathione-S-transferase GPx elutathione pervidase GP elutathione	atalase, FCx ro	ntal cortex, <i>G6PDH</i> glucose-6-ph transferase <i>GPv</i> obitathione new	osphate dehydrogenase, oxidase <i>GR</i> alutathione

Abbreviations: $\uparrow$ increase, $\downarrow$ decrease, $\emptyset$ no effect, D-Amph D-amphetamine, CAT catalase, FCx rontal cortex, G6PDH glucose-6-phosphate dehydrogenase,
GGTP y-glutamyltranspeptidase, GSH glutathione, GSSH reduced glutathione, GST glutathione-S-transferase, GPx glutathione peroxidase, GR glutathione
reductase, HIP hippocampus, HYP hypothalamus, LPO lipoperoxides, MDA malondialdehyde, MDMA 3,4-methylenedioxy-N-methylamphetamine, METH
methamphetamine, NAc nucleus accumbens, 3-NT 3-nitrotyrosine, PFCx medial prefrontal cortex, PC protein carbonyls, SOD superoxide dismutase, STR
striatum

Model treatment	Animal	Dose and treatment regimen	Biomarker	Change (brain structure)	Reference
Chronic passive administration	Rat	20 mg/kg ( <i>i.p.</i> ) for 10 days	GPx	†( FCx, STR), Ø(CER)	Dietrich et al. (2005)
			SOD	↑(FCx), Ø(STR)	
			LP	†(FCx, STR), Ø(CER)	
Chronic passive	Rat	15 mg/kg (i.p.)	GSH	↓(HIP)	Muriach
administration		for 20 days	GPx	↓(HIP)	et al. (2010)
Chronic passive administration	Rat	15 mg/kg ( <i>i.p.</i> ), 2 x in 1 and 7 days; 30 mg/kg ( <i>i.p.</i> ) for 2–6 days	GSH	↓(NAc)	Uys et al. (2011)
Subchronic passive administration	Prenatal rat	20 mg/kg ( <i>i.p.</i> ) for 4 days	MDA	↑(FCx, HIP)	Bashkatova et al. (2006)
Chronic self-	Rat	15-18 mg/kg/day	SOD	↑(STR, FCx, HIP)	Pomierny-
administration and extinction model		( <i>i.v.</i> ) +10-day extinction	MDA	$\downarrow$ (STR, FCx, HIP)	Chamiolo et al. (2012)
			SOD	$\uparrow$ (STR, FCx, HIP)	
			MDA	$\uparrow$ (STR, FCx, HIP)	
Acute administration	Rat	30 mg/kg (i.p.)	MDA	↑(PFCx, NAc)	Numa et al. (2008)
Acute administration	Mouse	10 mg/kg ( <i>i.p.</i> ) 30 mg/kg ( <i>i.p.</i> ) 90 mg/kg ( <i>i.p.</i> )	CAT	$\frac{\downarrow(\text{STR}), \emptyset(\text{PFCx})}{\downarrow(\text{STR}), \emptyset(\text{PFCx})} \\ \frac{\downarrow(\text{STR}), \emptyset(\text{PFCx})}{\downarrow(\text{STR}), \downarrow(\text{PFCx})}$	Macedo et al. (2005)
Acute	Rat	90 mg/kg (i.p.)	GSH	↑(PFCx, STR)	Macedo et al.
administration			MDA	↑(PFCx, STR)	(2010)

 Table 2
 Preclinical evidence of oxidative stress after cocaine exposure

**Abbreviations:**  $\uparrow$  increase,  $\downarrow$  decrease,  $\emptyset$  no effect, *CER* cerebellum, *FCx* frontal cortex, *GPx* glutathione peroxidase, *GSH* glutathione, *HIP* hippocampus, *LP* lipid peroxidation, *MDA* malondialdehyde, *NAc* nucleus accumbens, *PFCx* prefrontal cortex, *SOD* superoxide dismutase, *STR* striatum

(Krasnova et al. 2001) in animals and in humans, as hyperthermia – particularly if ambient temperature is high – promotes greater RS formation and associated neurotoxicity (Green et al. 2003; Quinton and Yamamoto 2006; Parrott 2012).

Cocaine likewise was found to promote oxidative stress in the brain of experimental animals (see Table 2), and it was recorded as a significant elevation in SOD, glutathione peroxidase (GPx), and CAT activity in the rat cortex and striatum (Dietrich et al. 2005; Macedo et al. 2005, 2010); however, changes in brain CAT were not validated in other studies (Macedo et al. 2005; Muriach et al. 2010). In the hippocampus GSH concentration and GPx activity were reduced in repeatedly cocaine-treated rats; such changes were not evident in the frontal cortex (Muriach et al. 2010). Cocaine intoxication led also to an increase in lipid peroxidation in several brain structures linked to dopamine synthesis and release (Dietrich et al. 2005; Bashkatova et al. 2006). A very recent report indicates that voluntary cocaine intake (by *i.v.* drug self-administration) evokes a significant reduction in MDA level together with increases in SOD activity in the hippocampus and frontal cortex, while an enhancement in the SOD activity was not seen in yoked cocaine animals (Pomierny-Chamiolo et al. 2012). As suggested, enhanced RS production and oxidative cell status may occur via cocaine or its oxidized metabolites (e.g., norcocaine derivatives; Kovacic 2005).

As stated above, psychostimulants prompt an increase in RS level via enhanced synaptic dopamine concentration or by its oxidation by nonenzymatic or enzymatic pathways (see above). In the mechanisms of drug dependence and neurotoxicity, glutamatergic neurotransmission is strongly involved. In cocaine addiction an increased level of extracellular glutamate concentration in dopamine-innervated areas (Cunha-Oliveira et al. 2008) could be associated with excitotoxic processes through increased activation of NMDA receptors and subsequent increase in intracellular Ca<sup>2+</sup> level (Rego and Oliveira 2003), leading to the formation of mitochondrial RS and lipid peroxidation (Maciel et al. 2001).

Antioxidant status after chronic exposure may be different than effects of acute exposure due to maladaptations seen in neurotransmitter levels and in antioxidant defense mechanisms that maintain cellular homeostasis (see Table 1 for specific changes in a brain region, drug type, and dosage regimen).

### Manifestation of Apoptosis

Several preclinical studies have reported apoptotic features after chronic or even acute exposure to psychostimulants, especially amphetamine derivatives (Cunha-Oliveira et al. 2008). For example, repeated amphetamine administration in mice  $(4 \times 10 \text{ mg/kg}, \text{ every } 2 \text{ h})$  caused an increase in p53 protein levels in medium spiny striatal projection neurons with a secondary increase in Bax (proapoptotic gen) and decrease in Bcl-2 (anti-death gene) levels in mouse striatum (Krasnova et al. 2005). Neurotoxicity due to activation of apoptotic pathways has been shown with acute drug treatments. Thus, acute methamphetamine (40 mg/kg, *i.p.*) resulted in an activation proapoptotic Bcl-2 family genes (Bax, Bad and Bid) in mice that was accompanied by a decrease in anti-death genes Bcl-2 and Bcl-X in the frontal cortex (Jayanthi et al. 2001). The same methamphetamine treatment resulted in the release of AIF and smac/DIABLO from the mitochondria in mouse striatum (Jayanthi et al. 2004), followed by activation of the caspase cascade (caspase 9 and 3) (Deng et al. 2002; Jayanthi et al. 2004). Similarly to methamphetamine, acute MDMA exposure caused upregulation of Bax and downregulation of Bcl-2 in the rat hippocampus and cerebellum and respective changes in genes relating to those proteins (Soleimani Asl et al. 2011, 2012). The latter findings suggest that MDMA neurotoxicity can be linked to stimulation of the intrinsic apoptosis pathway. Of note, apoptotic biomarkers were evident also in rat cortical neurons following MDMA (Capela et al. 2006).

In addition to the above mitochondrial apoptosis pathway, acute high-dose (40 mg/kg, *i.p.*) methamphetamine was found to activate cell death via

dysregulation of calcium homeostasis in mice (Jayanthi et al. 2004), initiating calcium-dependent apoptosis via the permeabilization of the outer mitochondrial membrane (Kroemer et al. 2007). Furthermore, acute methamphetamine injection with 20–40 mg/kg to rodents caused activation of calpain – a calcium responsive cytosolic cysteine protease (Jayanthi et al. 2004; Samantaray et al. 2006). In line with the latter observations, proteolysis of the cytoskeletal scaffolding protein spectrin, consequent to calpain activation, has been shown in many studies in rat brain limbic and subcortical structures following methamphetamine exposure (Warren et al. 2005; Staszewski and Yamamoto 2006). Finally, toxic doses of acute methamphetamine (40 mg/kg, *i.p.*) have been shown, in rat striatum, to induce the transcription factors AP-1, Egr, and Nur77 (Jayanthi et al. 2001). Altogether, the literature findings indicate several factors in the amphetamine-related cell death in rodent brain, including activation of either the mitochondria-dependent apoptotic pathway or activation of calpain, or Fas death pathway, or p53 death pathway.

On the other hand, morphological changes of apoptosis induced by cocaine have not been evident (Cunha-Oliveira et al. 2008). As recently reported, chronic cocaine, including its 3-day withdrawal, altered neither Fas–FADD receptor complex, cytochrome c, caspase 3, AIF, and PARP-1 cleavage nor associated signaling in the cerebral cortex. However, some discrete changes (a 28 % decrease in the content of antiapoptotic truncated isoform of dopamine- and cAMP-regulated phosphoprotein of 32 kDa) in rat brain cortex appeared (Alvaro-Bartolomé et al. 2011).

#### Manifestations of Genomic and Epigenomic Actions

The regulation of changes in gene expression evoked by psychostimulant drugs of abuse is an area of intensive research (Cadet and Krasnova 2009; Piechota et al. 2012). Altered gene expression [e.g., brain-derived neurotrophic factor (BDNF), G-protein signaling 3] in drug addiction is reported in the prefrontal cortex, ventral tegmental area, and striatum (including the nucleus accumbens) (Wong et al. 2011). Another long-term molecular effect relates to upregulation of the cAMP pathway in the nucleus accumbens and dorsal striatum – partially mediated by CREB. Dopamine  $D_1$  receptor stimulation during chronic cocaine administration induces activation of transcription factor CREB, which has many gene targets including BDNF, c-Fos, adenylyl cyclase, corticotropin-releasing factor, dynorphin, and enkephalins (Cunha-Oliveira et al. 2008). A recent study in mice, using wholegenome microarray profiling to evaluate the detailed time course of transcriptional alterations, identified 27 genes (i.e., circadian genes Per1, Per2, and Nr1d1; the glucocorticoid-dependent genes Fkbp5, Sult1a1, and Plin4; the drug-regulated group of transcripts enriched in the nucleus accumbens Pdyn, Cartpt, and Rgs2; as well as new genes Fam40b and Inmt) under regulation by chronic methamphetamine administration (Piechota et al. 2012). As suggested, the above transient gene expression alterations during drug treatment – seen also in the early period of withdrawal - may be contributors to persistent neuroplastic alterations responsible for the development of drug addiction.

A number of findings suggest that epigenetic changes may mediate neuroplasticity in psychostimulant addiction (Kumar et al. 2005; Levine et al. 2005; Romieu et al. 2008; Im et al. 2010; Maze et al. 2010). A recent study has demonstrated that amphetamine induced phosphorylation of methyl CpG binging protein (MeCP2) at Ser421 in mouse accumbal areas, leading to increases in protein activity and enhanced repressive function (Deng et al. 2010). Findings by Renthal and Nestler 2008 show that acute exposure to amphetamine results in  $\Delta$ FosB binding to the *c-fos* promoter and in repression of its expression through recruitment of HDAC1 and histone methyltransferase (HMT) around the promoter site, leading to downregulation of the gene expression. The same authors showed that chronic amphetamine abuse increases histone H3K9me2 on the *c-fos* promoter, further substantiating its transcriptional silencing (Renthal and Nestler 2008). As suggested, interaction between  $\Delta$ FosB and *c-fos* may serve as a homeostatic mechanism for the regulation of sensitivity to repeated drug treatment. Other studies indicated global increase in histone H4 acetylation and specific fosB promoterassociated H4 hyperacetylation in mouse striatum after repeated amphetamine treatment (Shen et al. 2008). Moreover, co-treatment of amphetamine and HDAC inhibitors (butyric acid and valproic acid) had additive effects on the increase in histone H4 acetylation (Shen et al. 2008; Adachi and Monteggia 2009). The latter study seems to support the notion that amphetamine functions as an HDAC inhibitor, suggesting that HDACs could be used therapeutically for the modulation of psychostimulant action. Acute amphetamine administration provokes an increase in the number of neurons positive for phosphorylation of histone H3 in serine 10 (pH3S10) in a few telencephalic and diencephalic rat brain regions (Rottlant and Armario 2012).

Acute methamphetamine exposure decreased DNMT1 and DNMT2 in the rat hippocampus (Numachi et al. 2004), while bidirectional changes for DNMT1 were observed in the nucleus accumbens in two rat strains (Numachi et al. 2007) following subchronic methamphetamine exposure. The fact that Lewis rats are more susceptible than Fisher rats to methamphetamine sensitization may be reflective of different DNA methylation patterns (Numachi et al. 2007; Kovatsi et al. 2011). Interestingly, a single methamphetamine injection (20 mg/kg) induced timedependent modifications in the acetylation of histone H3 and H4 as well as increases in AFT2 (enzymes that have HAT activity for histone H4) and HDAC2 expression in rat nucleus accumbens (Martin et al. 2012). Martin and coworkers demonstrated methamphetamine-induced increases of histone H4K5 and H4K8 acetylation as well as decreases in the acetylation of histone H3K9, H3K16, and H3K18. These results suggest that the psychostimulant effect mediated by increased AFT2 and HDAC2 expression may relate to effects on histone acetylation and hypoacetylation. The most recent study has indicated that repeated methamphetamine injections significantly decreased acetylation of histone H4 at lysine 5 (H4K5), lysine 12 (H4K12), and lysine 16 (H4K16) in rat dorsal striatum, at *GluA1* and *GluA2* DNA sequences encoded  $\alpha$ -amino-3-hydroxy-5-methyl-4isoxazolepropionic acid (AMPA) receptor subunits (Cadet and Jayanthi 2013). Because these changes were seen with the enrichment of methyl CpG binding protein 2 (MeCP2) at the promoters of both *GluA1* and *GluA2*, the results indicate that methamphetamine induced downregulation of *GluA1* and *GluA2* mRNA levels and did affect glutamatergic function.

Some limited data indicate amphetamines affect expression patterns of microRNAs (miRNA)s. Thus, Lippi and colleagues (2011) have shown that chronic treatment with amphetamine increases the level of *miR-29a/b* in mouse hippocampus. The *miR-29a/b* family directly targets the transcript of *Arpc3*, a subunit of the ARP2/3 actin nucleation complex involved in dendritic spine morphogenesis (Lippi et al. 2011). Current bioinformatic analysis has revealed that amphetamine increases levels of *miR-181a* in a mouse model of chronic drug treatment (Saba et al. 2012), linking epigenetic modification with synaptic plasticity. In fact, *miR-181a* is a negative posttranscriptional regulator of GluA2 expression, and thus it would be an essential regulator of mammalian AMPA-type glutamate receptors, a key determinant of synaptic plasticity.

There are some links between DNA methylation and cocaine addiction. First, cocaine self-administration increased MeCP2 expression in rat ventral and dorsal striatum (Cassel et al. 2006), while lentiviral knockdown of MeCP2 in dorsal striatum decreased drug intake under extended drug conditions (Im et al. 2010). Repeated cocaine administration altered DNMT3a, an enzyme linked to CpG methylation, expression in the mouse nucleus accumbens, with decreases seen during very early phases of withdrawal and sustained increases seen at later time points (LaPlant et al. 2010). Of note, prenatal cocaine exposure alters DNA methylation and gene expression in hippocampal neurons of neonatal and adolescent mice (Novikova et al. 2008).

A few reports point also to interaction between histone acetylation and cocaine. Thus, acute or chronic exposure to this psychostimulant increased cellular levels of H3 and H4 acetylation in the nucleus accumbens (Kumar et al. 2005; McQuown and Wood 2010), while repeated passive cocaine treatment downregulated dimethylation of histone 3 lysine 9 (H3K9me2), a marker of heterochromatin in the same brain area (Maze et al. 2010); this later effect paralleled a decrease in the total amount of heterochromatin and an increase in the volume of these nuclei. Increased H3 acetylation was also observed during cocaine self-administration and conditional place preference, while pharmacological inhibition of sirtuins (their induction is associated with H3 acetylation) reduced cocaine rewarding effects seen in conditioned place preference and self-administration (Renthal et al. 2009).

Findings from the past 3 years indicate that chronic cocaine regulates numerous miRNAs in mouse striatum (Schaefer et al. 2010; Eipper-Mains et al. 2011) as well miRNA-linked proteins that are strongly linked to drug addiction (Li and van der Vaart 2011). For example, cocaine self-administration in rats increased *miR-212* expression in the striatum that was suggested to be – at least partially – responsible for cocaine overconsumption through CREB, MeCP2, and BDNF signaling (Im et al. 2010). Another report identified induction of *miR-212* and *miR-132* in the dorsal striatum after 7 days of cocaine self-administration in rats (Hollander et al. 2010). On the other hand, chronic passive cocaine administration decreased *miR-124* 

and increased *miR-181a* in rat brain (Chandrasekar and Dreyer 2009). *miR-124* overexpression in the nucleus accumbens reduced cocaine place conditioning, whereas overexpression of *miR-181a* had the opposite effect (Chandrasekar and Dreyer 2011). Recently, it was found that cocaine mediates the changes of miR-8 family expression in brain and basing on the biostatistical analyses, a suggestion of novel mechanisms for drug-induced alterations in the neuronal cytoskeletal and synaptic structure has been raised (Eipper-Mains et al. 2011). Interestingly, Schaefer et al. (2010) identified an overlapping subset of cocaine-induced miRNAs (*miR-324* and *miR-369*) in dopamine D<sub>2</sub> receptor-expressing neurons of the nucleus accumbens together with reduction in cocaine self-administration.

# 4.2.2 Opiates

### **Metabolic Changes**

An early report on a single dose of morphine (20 mg/kg) demonstrated increases in glucose concentration (42 %) in rat brain (Miller et al. 1972). The same enhancement in local cerebral metabolic rate for glucose was shown for morphine repeated treatment in 95 brain regions of rats; the most significant changes were seen in the nucleus accumbens and prefrontal cortex (Kraus et al. 1997).

#### **Morphological Changes**

Chronic treatment of morphine (from 20 mg/kg to 140 mg/kg twice a day over 14 days at escalating doses) in rats, followed by drug withdrawal, resulted in a dramatic reduction of spine density in second-order dendritic trunks in the accumbens shell (Spiga et al. 2005). Chu et al. (2008) using the same treatment protocol with morphine showed that rough endoplasmic reticulum swelled, membrane configuration of the nucleus and mitochondria were blurred, and structure of myelin sheath changed in the rat ventral tegmental area. These findings support the linkage between opioid-induced abuse and structural functional plasticity of dendritic spines in the brain.

### Glia and Microglia Activation

Opioid-induced proinflammatory glial activation (increases in microglial and astrocytic activation markers and release of proinflammatory cytokinesis) was suggested to be involved in the development of opioid dependence and reward (Watkins et al. 2005; Hutchinson et al. 2007). Recently, the glial markers have been implicated also in drug-induced craving. Thus, Lin et al. (2011) showed that repeated exposures to morphine decreased glial fibrillary acidic protein in rat amygdala tissue in the stage of extinction. Another report indicates that heroin self-administration and withdrawal regulate in a time-dependent manner tegmental and accumbal glial cell line-derived neurotrophic factor mRNA expression (Airavaara et al. 2011).

## Manifestation of Oxidative Stress

There is evidence that repeated opioid treatments induce oxidative stress in the brain (see Table 3). Heroin injected continuously in increasing dosage

Model treatment	Animal	Dose and treatment regimen	Biomarker	Change (brain structure)	Reference
Morphine, chronic passive administration	Mouse	5 mg/kg ( <i>s.c.</i> ) for 7 days	GSH	↓Brain homogenate	Abdel-Zaher et al. (2010)
			GPx	↓Brain homogenate	
			MDA	†Brain homogenate	
Tramadol, chronic passive administration	Mouse	50 mg/kg ( <i>s.c.</i> ) for 7 and 15 days	GSH	↓Brain homogenate	Abdel-Zaher et al. (2011)
			GSH-Px	↓Brain homogenate	
			MDA	†Brain homogenate	

 Table 3
 Preclinical evidence of oxidative stress after opioid exposure

**Abbreviations:**  $\uparrow$  increase,  $\downarrow$  decrease, GPx glutathione peroxidase, GSH gluthatione, MDA malondialdehyde

(1–35 mg/kg) to mice caused increasingly greater numbers of DNA-damaged cells, protein carbonyl groups, and MDA levels (Qiusheng et al. 2005). Prooxidant status was also marked by decreased activity of antioxidant enzymes (SOD, CAT, GSH) (Qiusheng et al. 2005). It should be stressed that even a single morphine dose can produce oxidative damage in the brain (e.g., decreased GSH levels); however, its manifestation depends on the dose of morphine and age of the animal. Another opioid, tramadol at a 50 mg/kg (*i.p.*) dose, decreased nonenzymatic antioxidants such as intracellular GSH and enzymatic antioxidants such as GPx in mouse brain; repeated tramadol treatments resulted in an increase in the MDA level in the brain (Abdel-Zaher et al. 2011).

#### Manifestation of Apoptosis

Several studies performed on rodents have shown that sustained exposure to opioids produced cell death in many areas of the brain (Mao et al. 2002; Emeterio et al. 2006), together with increases in the brain expression of proapoptotic factors belonging to both the mitochondrial and the Fas-receptor apoptotic pathways (Boronat et al. 2001; Mao et al. 2002; Emeterio et al. 2006; Tramullas et al. 2007). For example, prolonged (34 days) heroin (5 mg/kg, *s.c.*) administration to mice evoked upregulation of proapoptotic proteins such as Fas, FasL, and Bad, in the cortex and hippocampus, indicating the activation of both the death receptor and the mitochondrial apoptotic pathways. In parallel, the presence of TUNEL (terminal deoxyribonucleotidyl transferase-mediated dUTP nick and labeling method)-positive cells scattered throughout the brain evidenced another indicator of apoptosis linked to DNA fragmentation (Tramullas et al. 2008). The signs of apoptosis were reported even after a single injection of heroin (5 mg/kg), which increased the expression level of Bad in the cortex and FasL in the hippocampus.

These biochemical indices in heroin-treated mice were accompanied by impaired cognitive ability in the Morris water maze and cognitive flexibility-related performance, raising the possibility that chronic exposure to opioid drugs in the brain would interfere with learning and memory through a neurotoxic effect related to activation of apoptotic pathways. Interestingly, heroin-induced apoptosis (caspase 2 and 9 activation) in neuronal-like PC12 cells was associated by increased mitochondrial membrane permeability and decreased mitochondrial potential (Oliveira et al. 2002, 2003).

Similarly to heroin, other opioid drugs can lead to apoptosis in neuronal cells. Thus, tramadol and morphine given to rats in progressive doses of (20, 40, and 80 mg/kg/day of tramadol; 4, 8, and 12 mg/kg/day of morphine) significantly enhanced the presence of red neurons (a histologic marker of apoptosis) as compared with control group (Atici et al. 2004). Chronic morphine or tramadol use in increasing doses caused red neuron degeneration in the rat brain, which probably contributes to cerebral dysfunction. Interestingly, the numbers of red neurons were significantly higher in the temporal and occipital regions in the tramadol group as compared with the morphine group, and these findings are of importance when prolonged opioid treatment is necessary (Atici et al. 2004).

### Manifestation of Genetic and Epigenetic Actions

Genetic and epigenetic changes are another area of molecular interactions that could explain the mechanism of opioid neurotoxicity, addiction, and abstinence syndrome. In numerous studies altered adenyl cyclase activity and enhanced intracellular levels of cAMP have been observed during manifestations of opioid withdrawal syndromes (Sharma et al. 1977; Law et al. 1982; Koob and Bloom 1988; Piñeyro and Archer-Lahlou 2007). Recently reported by Piechota et al. (2012), chronic treatment with heroin regulates several genes in mouse striatum, with the most substantial impact for circadian genes (Per1, Per2, and Nr1d1) and glucocorticoid-dependent genes (Fkbp5, Sult1a1, and Plin4), a drug-regulated group of transcripts (Pdyn, Cartpt, and Rgs2, as well as Fam40b and Inmt). The appearance of opioid receptor downregulation and their desensitization were suggested to be the result of histone-deacetylase-mediated epigenetic control of genes (Hu et al. 2001; Hwang et al. 2007). Another paper by Zhou et al. (2006) revealed that morphine-withdrawal-related stress condition increased oprml expression in the rat hypothalamus and striatum as well as lateral hypothalamic orexin mRNA level; this finding supports the hypothesis that orexin activity plays a role in morphine-withdrawal-related behaviors. There is also a finding showing that morphine decreased acetylation of histones H3 and H4, trimethylation of lysine 4 of H3 (H3K4me3) at the promoters of KCNAB2 and KCNF1 genes (both encoded subunits of potassium channel), which in turn decreased their transcription in rat ventral tegmental area (Mazei-Robison et al. 2011). Parallel reduction of trimethylation of H4K4me3 at the promoter of GIRK3 gene (encoded potassium subunit channel) may suggest that morphine downregulates the K<sup>+</sup> channel via epigenetic changes and enhanced excitability of dopamine cell bodies within the mesencephalon.

There are implications for the significance of noncoding RNA, primarily miRNAs, in opioid-based addictive behaviors. Morphine has been implicated in the regulation of *miR-23b* and *let-7* expression that regulates posttranscriptional levels of mu opioid receptor (Wu et al. 2009; He et al. 2010). In the rat hippocampus, morphine activates ERK1/2 and decreases the expression level of *miR-190* (Zheng et al. 2010), which regulates the expression of neurogenic differentiation 1, thereby affecting the morphology of dendrites and adult neurogenesis. In addition, *miR-190* interacts with *pax6* gene-encoded transcription factor regulating neurogenesis (Zheng et al. 2012).

## 5 Conclusion

Psychostimulants and opioids evoke toxicity within the CNS, which in human research is detected after a particular cumulative drug exposure, while in preclinical studies neurotoxicities are recorded even after acute drug administration. Mechanisms leading to neurotoxicity linked to psychostimulants and opioids are due primarily to the action of the substance (e.g., enhancement of synaptic concentrations of dopamine) or by indirect effects (increases in extrasynaptic glutamate levels, toxic metabolites). Associated with the development of drug addiction, these mechanisms lead to the several pathological processes within the complex network of brain cell-to-cell interactions, such as alteration in neurotransmitter systems, dysregulation of energy metabolism, changes in structure of cellular membrane, apoptosis, generation of RS, glial/microglial cell overactivation, excitotoxicity and prominent deficits in brain morphology (neuronal loss, axonal damage), alterations of transcription factors, immediate early genes and epigenetic mechanisms, as well as neurocognitive and neuropsychological impairments (e.g., risk of Parkinson's disease). The extent of neurotoxic variations depends on drug type, drug dosage (chronic vs acute treatment), species (human vs nonhuman primates or rodents), external factors (environmental temperature), and brain region.

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# Molecular Mechanism and Effects of Clostridial Neurotoxins

# Bal Ram Singh, Raj Kumar, and Shuowei Cai

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#### Abstract

The genus Clostridia, gram-positive anaerobic bacteria, produce two of the deadliest toxins known to mankind: tetanus and botulinum. Toxicity of these molecules comes from their action on very specific substrates, SNARE proteins, in the neuronal cells. They cleave SNAREs, which are the crucial components of neuroexocytosis, causing inhibition of release of neurotransmitters at the synapses. Botulinum neurotoxin molecule has evolved itself to have high stability, specificity, and selectivity. It maneuvers and exploits itself in each step of intoxication and preserves its active structure to achieve an exquisite toxicity. The mechanisms of action and biological functions of clostridial neurotoxins

e-mail: bsingh@umassd.edu; rkumar@umassd.edu; scai@umassd.edu

B.R. Singh (🖂) • R. Kumar • S. Cai

National Botulinum Research Center and Department of Chemistry and Biochemistry, University of Massachusetts Dartmouth, Dartmouth, MA, USA

still have several unresolved questions. Several unique characteristics of these neurotoxins attract scientists to understand the mysteries of neurotoxin molecules. Despite high toxicity, botulinum neurotoxin has a wide range of medicinal applications. In the last few decades, toxin transformed itself from lifethreatening disease to a wonder drug. Therapeutic use of botulinum neurotoxin is based on three unique features: specificity, potency, and duration of action. Active research is underway to transform botulinum neurotoxins into the most versatile and useful drugs for neuromuscular disorders, and as cosmeceuticals. This chapter summarizes current knowledge on its structure and mechanism of intoxication. It also describes basic understanding of the evolution of this molecule to achieve this extremely high biological activity in terms of toxicity or therapeutics. Understanding the true nature and function of its dynamic structure still remains a work in progress.

## 1 Toxins

In general toxins can be small molecules, peptides, or proteins. These molecules are toxic because they are capable to evade protection mechanisms of the infected host cells and affect important cellular processes such as protein synthesis, inhibition of other cellular processes, and promotion of hemolysis. Toxins can be generally grouped according to their source (e.g., bacterial toxins), their purpose (e.g., phototoxins), and their site of action or affected organs (e.g., hepatotoxins, neurotoxins). Among these the most complex and deadly toxins are the bacterial toxins. Pathogenic bacteria use many ways to interact with target tissues or cells. Toxins produced by bacteria can act locally or at a distant location from the sites of infection, leading to various diseases in human and animals. Bacterial toxins are classified as two types: exotoxins, which are secreted by the bacteria in the environment, and endotoxins, which may be released after the lysis of bacterial cell wall. Clostridial neurotoxins produced by anaerobic Clostridium botulinum and *Clostridium tetani* are exotoxins, which can be produced in infected tissues, such as wounds or infant guts. These neurotoxins are the most toxic substances currently known to mankind. There are many unique features associated with these neurotoxins and also with the bacteria which produce them, all of which together play a role in the mechanism of pathogenicity, making this class of pathogens highly unique.

# 2 Clostridium Family

The genus *Clostridium* represents gram-positive, ubiquitous, anaerobic, and endospore-forming bacteria. Clostridium has about 100 species consisting of ubiquitous bacteria and pathogens. Members of this group of bacteria exhibit large straight or slightly curve-shaped bacterial cells. Members of this genus produce some of the

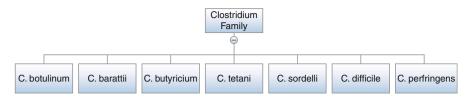


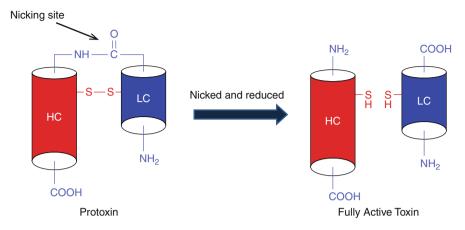
Fig. 1 Members of *Clostridium* family, known for producing toxins

most potent toxins discovered so far. Dormant cells are highly resistant to heat, desiccation, toxic chemicals, and detergents. Spores do not germinate normally unless they have suitable environment to grow. Most of clostridial species are saprophytes, but a few are pathogenic to humans and animals, for example, *C. difficile, C. botulinum, C. tetani*, and *C. perfringens* (Fig. 1). The pathogens are primarily saprophytes in nature. *C. tetani* and *C. botulinum* produce the most potent biological toxins known to mankind. In their active form they secrete exotoxins which cause various diseases, such as tetanus and botulism.

# 3 The Clostridium botulinum

*Clostridium botulinum*, which was first isolated from a contaminated ham and from the spleen of a patient by van Ermengem in Belgium in 1895 (Van Ermengem 1979; Foster 1993) grows anaerobically at 37 °C or 30 °C under pH 6.5–7.0 conditions. Most of C. botulinum as well as some isolates of C. butyricum and C. baratii can produce botulinum neurotoxins (BoNTs). The strains of C. botulinum are traditionally classified into seven serotypes, A to G, according to the antigenic properties of neurotoxins they produce. Generally, one strain of C. botulinum can only produce one type of neurotoxin, but several strains that produce more than one type of neurotoxin have been identified over the years. One type C strain (C6813) was found to produce the mosaic neurotoxin in which one-third of the C-terminal region was similar to that of type D neurotoxin (Moriishi et al. 1996). One type A strain, called type AB now, was shown to produce both type A and type B neurotoxins (Fujinaga et al. 1995). C. botulinum type G organism differs phenotypically from other C. botulinum types and is now designated as C. argentinense (Gimenez and Ciccarelli 1970; Campbell et al. 1993b). Clostridium butyricum and Clostridium *baratii* were found producing type E (BoNT/E) and type F (BoNT/F) neurotoxins, respectively (Fuji and Kimura 1993; Thompson et al. 1993).

BoNT is synthesized as a single and relatively inactive polypeptide chain, which undergoes a proteolytic cleavage at its nicking site to form a biologically active dichain structure linked by a disulfide bond. The proteolytic strains of *C. botulinum* produce endogenous proteases and secrete the neurotoxin as an active dichain form by cleaving it with endogenous protease, but non-proteolytic strains of *C. botulinum* require exogenous proteases, such as trypsin in the intestinal tract of hosts, to cleave the progenitor toxin to activate it (Call et al. 1995, Fig. 2).



**Fig. 2** Botulinum neurotoxin: BoNTs are produced as a single chain (progenitor or protoxin) which is processed exogenously or endogenously as a dichain. Reduction of disulfide bond makes fully active toxin

Strains of C. botulinum can also be divided into four phylogenetically and physiologically distinct groups (I to IV) based on biochemical properties and nucleic acid hybridization studies. Group I includes type A, proteolytic strains of types B and F. Group II includes type E, non-proteolytic strains of types B and F. Group III consists of types C and D, and group IV consists of type G (Campbell et al. 1993a). Rainey and Stackebrandt (1993) and Campbell et al. (1993b) used a very high level of 23S rRNA sequence similarity to show that groups I, II, III, and IV C. botulinum represent a single genetic unit phylogenetically remote from each other and that the marked genotypic variation within the species does not correlate with the neurotoxins produced. C. baratii that produces type F neurotoxin belongs to a new added group, group V, and *Clostridium butyricum* that produces type E neurotoxin belonged to the latest group, group VI (see Table 1; Chang 2011). The same conclusions were also drawn from 16S rRNA sequence similarities (Hutson et al. 1993). However, the genetic relationship inferred from the BoNT sequences differs markedly from above conclusions and it is clearly established now that all three types of BoNT/F (proteolytic, non-proteolytic, and C. baratii type F) are genealogically highly related.

The spores of proteolytic *C. botulinum* are heat resistant and do not grow at temperatures below 10 °C, but the spores of non-proteolytic *C. botulinum*, which are less heat resistant, can germinate relatively rapidly at temperatures in the range of 2–10 °C (Lund and Peck 1994). Type A *C. botulinum* has 0.39 h generation time and reaches its exponential growth after 5 h of anaerobic incubation of spores (Call et al. 1995). The bacterial growth curve of type A *C. botulinum* (strain Hall) shows a lag period of 4 h and log phase at 10 h before reaching the stationary phase (Roberts and Hobbs 1968; Billon et al. 1997; Shukla and Singh 1999, 2009; Stringer et al. 2009).

		Group I			Group II		Group III	∎d	Group IV	Group V	Group VI
Endogenous Protease Activity		Proteolytic			Non Proteolytic	tic	Non Proteolytic	teolytic	Proteolytic	Non Proteolytic	Non Proteolytic
Clostridial Neurotoxin Serotype	A	в	ш	в	ш	ш	υ	٥	g	ш	ш
Accumulated Subtypes so far	A1-A5	B1-B3 Bivalent B	F1-F5	B4,B5	F6	E1-E3 E6				F7	E4,E5
Belong to Type-I: Ha(+)/Orf-X(-)	A1, A5	B		В			υ	٥	σ		
Belong to Type-II: Ha(+)/Orf-X(-)	A1-A4		F1-F5		F6	ш				F7	ш
	SV2A, SV2B, SV2C	Syt I and II,	glycosylated SV2A,			SV2A, SV2B	Unknown	uwc	Syt I and II,		
Neuronal Receptors	GT1b, GD1a	GT1b, GD1a	GT1b, GD1a			GT1b, GT1a		ЪЕ	all ganglioside		
	GD1b	GD1b	GD1b								
Toxins Specific Substrate	SNAP-25		VAMP (also called Synaptobrevin)	tptobrevin)		SNAP-25	SNAP-25 Syntaxin		VAMP (also called Synaptobrevin)	aptobrevin)	SNAP-25
	Type E and F that co	me from differer	it groups all belong to r	leurotoxin ger	ne cluster typ	e II; <i>cnt</i> R is at	sent from all	type E and	Type E and F that come from different groups all belong to neurotoxin gene duster type II; cnt R is absent from all type E and cnt C is absent from all type G.	type G.)	
Harbored Neurotoxin genes	io.	Single and Bivalent	t						Single		
Bivalent Gene Types if contained	A	Ab, Af, Ba, Bf, A(B)	(8)					Hasn	Hasn't Found Yet		
Mean Genome Size		4.02 Mb			3.73 Mb		2.96 Mb	qly			
Mean % G + C of Genome		28.23%			27.45%		29.00%	%			
Neurotoxin Gene Site	Chro	Chromosome or Plasmid	smid	Chro	Chromosome or Plasmid	lasmid	acteriophage	hage	Plasmid	Transposon	Transposon
Non-Neurotoxigenic				-			C. noyyi	yyi	C. subterminate	0 1	
Equivalent clostridia		C. sporogenes		Las	nasn t given name yet	ne yet	C. hemolyticum	vticum	C. proteolyticus	iypical C. baratil	Iypical C. bulyricum
Growth Temperature Attribution		Mesophilic			Psychrotropic	ic	Mesophilic	hilic	Mesophilic	Mesophilic	Mesophilic
Shore Heat Resistance		$D_{121^{\circ}C} = 0.21 \text{ min}$	-		$D_{82.2^{\circ}C} = 2.4 \text{ min}$	nin	D <sub>104°C</sub> = 0.91 min	.91 min	D <sub>104°C</sub> = 1.1 min		D <sub>100°C</sub> < 0.1 min
	Hig	High Heat Resistance	ce	Mode	Moderate Heat Resistance	sistance	Moderate	ate	Moderate	Moderate	Moderate
	P	Foodborne Botulism	E				Avian Botulism	tulism	Toxin is toxic in vitro	Foodborne Botulism	Infant Botulism
Associated Botulism Types		Wound Botulism		Fo	Foodborne Botulism	nlism	Foxes Botulism	tulism	but hasn't found	Infant Botulism	Adult-Intestinal
	Infar	Infant/Intestinal Botulism	ism				Minks Botulism	tulism	case report yet	Adult-Intestinal	
Optimal Growth °C		37°C			25°C		40°C	0	37°C	30~45°C	30~37°C
Minimum Growth °C		10~12°C			2.5~3.0°C		15°C			10~15°C	12°C
Minimal pH for Growth		4.6			5		5.1				4.8
[NaCl%] Preventing Growth		10%			5%				6.50%		
Lipase Production		+			+		+		1	I	1
Glucose Fermentation		+			+		+		I	+	+
Fructose Fermentation		-/+			+		-/+		I	+	+
Maltose Fermentation		-/+			+		-/+		1	+	+
Sucrose Fermentation		ı			I		1		1	+	+
	Source: Data from Smit	th & Sugiyama (	Source: Data from Smith & Sugiyama (1988), Lund and Peck (2000), Stringer and Peck (2008), Poulain et al. (2008)	2000), Stringe	er and Peck	2008), Poulair	et al. (2008)				
	means all strians ; +/- r	neans some stra	means all strians ; +/- means some strains are positive while others are negative; - means all strains negative.	thers are neg	jative; – mea	ns all strains n	egative.				
Remark	Group II could be separ	rated into 2 majo	Group II could be separated into 2 major groups in the future, one containing E the other containing B & F.	one containing	g E the other	containing B 8	Ш				
	The genome of a phage contains BoNT/C gene cluster has bee	e contains BoNT	The genome of a phage contains BoNT/C gene cluster has been sequenced. It is a linear dsDNA of 186 kb with a low %G+C (26.2%)	en sequence	d. It is a linea	r dsDNA of 18	5 kb with a lo	w %G+C (26	6.2%)		

The studies with a monoclonal antitoxin conjugated with colloidal gold demonstrated that the type A BoNT is synthesized in the cell cytoplasm at late-exponential phase (17 h) and reaches its maximum concentration at stationary phase (24–25 h). The BoNT is translocated across the cytoplasmic membrane and then exported to the extracellular medium via cell wall exfoliation. The greatest proportion of BoNT is located within the cell wall at 45 h (Call et al. 1995).

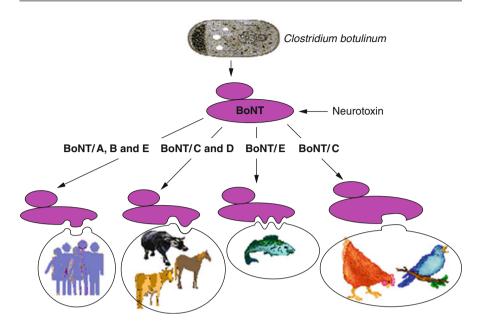
The spores of C. botulinum are widely distributed in soil, mud, and surface waters over much of the globe, but different types of C. botulinum prevail in different geographical locations. In the western United States, Brazil, Argentina, and China, type A spores predominate, while proteolytic type B spores tend to predominate in soils of the eastern United States and non-proteolytic type B in Britain and Europe. Type E is often associated with freshwater and marine sediments, and northern regions such as northern Canada and Japan (McClure et al. 1994). Initially, the strains of C. botulinum type A were divided into two groups, infant type A and type A, based on the cases of infant botulism in Japan and the classical food-borne botulism (Sakaguchi et al. 1990). But several studies (Collins et al. 1994; Cordoba and Collins 1995; Aranda et al. 1997) have revealed that the strains of C. botulinum type A could be divided into two groups (subtypes), A1 and A2, based on different geographical locations isolated and did not correlate with their origins as the cases of both infant botulism and food-borne botulism. Strains isolated from infant botulism in Japan and several strains isolated from classical food-borne botulism in United Kingdom and in Mauritius fell into group A2, while all strains examined from the USA and two strains from UK, from both infant and food-borne botulism fell, into group A1. Group A1 is typical C. botulinum type A containing hemagglutinin genes and shows hemagglutinin activity. Several group A1 strains contain an unexpressed type B neurotoxin gene (Ab type). Group A2 lacks the hemagglutinin gene and possesses no hemagglutinin activity. Along with the increasing number of sequenced type A strains, we can now identify more subtypes based on the criteria of sequence variations between 2.6 % and 32 % at the amino acid level, while compared to up to 70 % between different main serotypes of neurotoxin (Marks 2004; Smith et al. 2005; Arndt et al. 2006). Currently there are five subtypes of type A neurotoxin (termed A1, A2, A3, A4, A5), five type B subtypes, six type E subtypes (Arndt et al. 2006; Chen et al. 2007a; Hill et al. 2007; Smith and Hill 2007; Carter et al. 2009), and seven type F subtypes (Raphael et al. 2010) have been classified (see details in Table 1, Chang 2011). Interestingly all five type A subtypes belong to the same group (group I), while the other subtypes belong to more than one groups. Overall, the subtypes A1  $\sim$  A5, B1  $\sim$  B3, and F1  $\sim$  F5 belong to group I. The subtypes B4  $\sim$  B5, F6, E1  $\sim$  E3, and E6 belong to group II. The subtypes E4  $\sim$  E5 belong to group VI, and only subtype F7 belongs to group V. The existence of these subtypes is a manifestation of host adaption and divergent evolution of neurotoxin gene, and it will be important more and more in the development of neurotoxin antiserum and antidotes.

## 3.1 Botulism

Historically, botulism – derived from botulinus meaning sausage – was discovered long before the name appeared in the literature. Earliest reference to possible botulism-related disease is found in an edict by Emperor Leo VI of Byzantium (886–911) in which manufacturing of blood sausages was forbidden (Erbguth 2004). However, the first systematic study of "sausage poisoning" was carried out in 1815 by a health officer in the town of Herrenberg of Württemberg (Germany) by the name of J. G. Steinbuch and later independently by Justinus Kerner, another medical officer in the town of Welzheim in Württemberg (Erbguth 2004). Kerner's work was published in journals and monographs in early 1800s. In 1822, Kerner described his observations and experiments with sausage toxin, also referred as fat poison, in a monograph. Kerner wrote "The nerve conduction is brought by the toxin into a condition in which its influence on the chemical process of life is interrupted. The capacity of nerve conduction is interrupted by the toxin in the same way as in an electrical conductor by rust." Interestingly, Kerner had also hypothesized on the idea of sausage toxin being used as a therapeutic agent (Erbguth 1996, 1998).

Botulism is a severe disease characterized by flaccid paralysis and caused by botulism neurotoxins, extremely potent food poisons with a mouse lethal dose of 0.1 ng/kg (Montecucco and Schiavo 1993). Three types of botulism have been observed, so far. Classical food-borne botulism is caused by ingestion of food where the bacteria have produced the neurotoxin. Infant botulism result due to the production of neurotoxin in the intestine after germination and outgrowth of ingested spores. Wound botulism maybe due to the outgrowth of bacteria in the wound or due to unknown origin (Cordoba and Collins 1995; Hatheway and Ferreira 1996). Classical food-borne botulism in humans is caused mainly by C. botulinum types A, B, E, and rarely by type F neurotoxins. Classical foodborne botulism in animals is always caused by C. botulinum types C1 and D neurotoxins (Fach et al. 1993; Cordoba and Collins 1995). In contrast, the causative agents for infant botulism can be C. botulinum types A or B neurotoxins or Clostridium butyricum type E and Clostridium baratii type F neurotoxins. The wound botulism is rare in humans, and the causative agents are either type A or type B from group I (Hatheway 1993), but is increasingly significantly in recent years among drug users (Merrison et al. 2002). Neurotoxins can also be used in the therapy of a variety of human muscle disorders such as strabismus, hemifacial spasm, and blepherospasm (Schantz and Johnson 1992; Eleopra et al. 2004).

Epidemiologically, different serotypes of BoNT are known to selectively cause the disease in different animals. For example, BoNT/C is mostly associated with avian botulism; BoNT/C and BoNT/D with cattle, sheep, and horse; and BoNT/A, BoNT/B, and BoNT/E are associated with human, cattle, and horse botulism, while BoNT/E is particularly associated with fish botulism (Huss and Eskildsen 1974; Burns and Williams 1975; Ala-Huikku et al. 1977; Fernandez et al. 1989; Baldassi et al. 1991; Haagsma 1991; Jeffrey et al. 1994; Ashie et al. 1996;



**Fig. 3** Schematic representation of different binding topography of various serotypes of BoNT for their interactions with unique protein receptors in different species (Adapted from Singh 1999)

Ortolani et al. 1997). It is believed that botulism by a selective serotype of BoNT in a given animal is in part related to the presence of a high-affinity receptor for that serotype in the host animal system (Fig. 3). The true case number of food-borne botulism is probably underreported in the world. First, botulism is not a reportable disease in all countries (Therre 1999), and secondly the efficiency of investigating potential outbreaks varies from country to country (Peck 2009). It is estimated that the botulism case reporting efficiency in the USA is about 50 % only (Mead et al. 1999).

# 4 Clostridium Neurotoxins

Clostridial bacteria produce two of the deadliest neurotoxins known: tetanus and botulinum. Tetanus neurotoxin is produced by *C. tetani* and *C. botulinum* produces botulinum toxin. *C. tetani* produces a single toxic species, whereas strains of *Clostridium botulinum* produce at least seven serologically distinct neurotoxins: A–G. BoNTs are food poisons, whereas TeNT is not, and this has been attributed in part to the existence of the complex form of BoNT, consisting of toxin as well as neurotoxin-associated proteins (NAPs). There is no protein associated with tetanus toxin (Singh et al. 1995; Innoue et al. 1996; Minton 1995). NAPs are supposed to protect and enhance the activity of neurotoxin (Sharma and Singh 2004;

Kukreja and Singh 2007). However, Maksymowych et al. (1999) showed that there is no role of NAPs in enhancing the toxin activity. The extreme potency of these toxins lies in their specificity to the neuronal cells.

Clostridial neurotoxins belong to a group of bacterial toxins known as A-B type toxins consisting of at least two separate functional domains: one is responsible for binding and internalization and other is responsible for the intercellular action of the toxins. Both BoNT and TeNT are known to exert neuroparalytic effect in vertebrates (Humeau et al. 2000). Tetanus is structurally similar to BoNT but produces different symptoms upon intoxication. The action site is also different: BoNTs act on peripheral nervous system, while TeNTs act on central nervous system (Caleo and Schiavo 2009). Tetanus toxin induces spastic paralysis, while botulinum toxin induces flaccid paralysis. BoNTs inhibit the release of acetylcholine at the neuromuscular junctions (Simpson 1986, 2004), whereas TeNT inhibits the release of glycine and  $\gamma$ -amino butyric acid in the central nervous system (Schiavo et al. 2000). However, BoNT/A suppresses the release of acetylcholine, glutamate, noradrenaline, glycine, serotonin, GABA, and dopamine (Bozzi et al. 2006; Ashton and Dolly 1988; Verderio et al. 2004, 2007; Sutton et al. 2004; Costantin et al. 2005; Antonucci et al. 2008) in neuronal cells and synaptosomes. The differential site of BoNT and TeNT action may be due to differences in the binding affinity of the BoNT and TeNT for peripheral nerves vs. central nerves and also due to differential retrograde action of the toxins once internalized into nerve cells (Schiavo et al. 1993b, 1994b, 2000; Wang et al. 2012).

Clostridial neurotoxins are Zn-metalloprotease having very high specificity for its substrate: SNARE proteins involved in neurotransmitter release in the neuronal cells. The toxins selectively cleave the SNARE proteins; syntaxin, synaptobrevin (also known as vesicle-associated membrane protein or VAMP) and SNAP-25, but the final results are same – the hydrolysis of the target protein, blockage of neurotransmitter release, and a resultant paralysis. While BoNTs remain in the nerve terminal, the tetanus toxin is transported retrograde axonal transport into the cell body. Tetanus and botulinum toxins affect both somatic and autonomic nervous systems, but autonomic features are more common in tetanus.

All seven serotypes of BoNT are structurally similar. They are synthesized as a single-polypeptide chain of 150 kDa each which is processed by endogenous or exogenous proteases into an activated dichain form (Dekleva and Das Gupta 1989; Krieglstein et al. 1994; Singh and DasGupta 1989, Fig. 2). In each case, the dichain neurotoxin is composed of 100 kDa heavy chain (HC) and a 50 kDa light chain (LC), which is linked through a disulfide bond. Heavy chain consists of two domains of 50 kDa each: H<sub>C</sub> and H<sub>N</sub>; H<sub>C</sub> is binding domain and H<sub>N</sub> is the translocation domain. Structurally each toxin molecule has all three domains: binding domain (H<sub>C</sub>), translocation domain (H<sub>N</sub>), and catalytic domain (LC). Hc comprises two subdomains – a  $\beta$ -sheets domain (H<sub>CN</sub>) and a  $\beta$ -tree-foil (H<sub>CC</sub>).

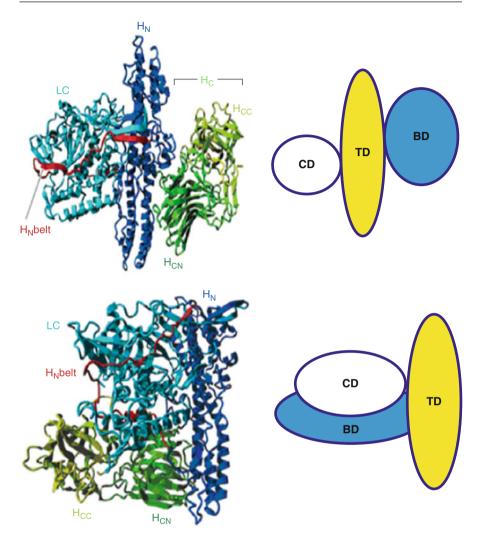
The three domains of BoNTs appear to be naturally designed for toxin delivery system, as binding domain (BD) acts a targeting unit, the translocation domain (TD)

can deliver cargo into a cell, and the catalytic domain acts as the cargo (Foster 2009; Bade et al. 2004; Chaddock and Marks 2006; Chaddock and Acharya 2011). The arrangement and the function of subunit in BoNTs are analogy to those in diphtheria toxin and anthrax toxin complex (Humeau et al. 2000; Monteccuco and Schiavo 1994; Montecucco et al. 1994; Montecucco and Schiavo 1995; Schiavo et al. 1994a, b; Singh 2000). In BoNT structure, in addition to the disulfide link between HC and LC, there is a belt, a loop from the HC that wraps around LC to hold the two chains together (Lacy and Stevens 1999; Chen et al. 2007b). The active site of LC is partially occluded by the belt in unreduced toxin (Simpson et al. 2004; Ahmed and Smith 2000), and the belt is predicted to be intermolecular chaperone for the light chain (Brunger et al. 2007). All the domains work together with high precision and accuracy. That is perhaps part of the reason why BoNTs are highly effective biologics, making them the most toxic substances.

The BD of HC binds to nerve cells, and after internalization of the BoNT through endocytosis, the TD forms a pore in the endosomal membrane, through which the LC domain crosses into the cytosol. This leads to intracellular blocking of neurotransmitters release by proteolysis of one of the three proteins of SNARE complex by Zn-endopeptidase activity of the LC. Isolated LC cleaves SNARE protein but it cannot readily internalize on its own, and therefore remain nontoxic. Also, isolated  $H_C$  or  $H_N$  cannot cleave SNARE proteins, so remain nontoxic. So only when three domains come together they create most the toxic substance known. It is notable that this unique assembly allows us to explore the possibility of combinatorial profiling of multidomain assemblies, for example, LC-TD for retargeting LC function (Darios et al. 2010).

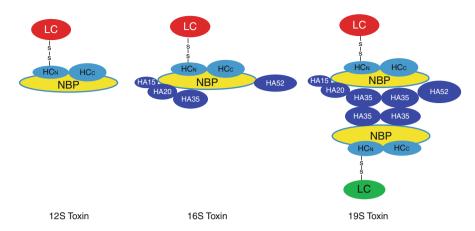
The crystal structure of BoNT/A (PDB: 3BTA), BoNT/B (PDB: 1EPW), and BoNT/E (PDB: 3FFZ) supports the hypothesis of modular assembly of BoNT (Fig. 4). Crystal structure of TeNT is not available. BoNT/A and BoNT/B show similar structural organization, whereas BoNT/E has some significant differences. Nevertheless, BoNT/E also has all the three domains, and while all other serotypes are made of two chains endogenously, BoNT/E is produced entirely as a single-chain protein. However, BoNT/B and BoNT/F are also produced as a single chain by non-proteolytic strains of the Clostridium botulinum (East et al. 1998). All the three domains are working together to achieve intoxication and act as a chaperone for the others (Brunger et al. 2007; Montal 2010). BoNT's functional complexities emerge from its modular design and tight interplay of individual component actions. Above-described design helps BoNT in using a very vital neuronal cell process, endocytosis, to reach to its targets. Exquisite toxicity of BoNTs is achieved by its unique structure which this molecule exploits very well. BoNT is considered an astonishing nanomachine that unites recognition, trafficking, unfolding, translocation, refolding, and catalysis in a single entity.

As stated earlier, TeNT is also synthesized by bacteria as a single-chain polypeptide, and because of the presence of bacterial protease, the toxin is isolated in a nicked form like botulinum toxin. Like A-B toxin, both separated heavy



**Fig. 4** Crystal structure of BoNT/A (3BTA) and BoNT/E (3FFZ). Hc is depicted in *green*,  $H_N$  is in *blue*, and LC is in *gray* color. Hc belt is depicted in *red*. BoNT/B (1EPW) is similar to BoNT/A. In case of BoNT/E LC (CD) and HC (BD) are on the same side of translocation domain (*TD*), whereas in BoNT/A CD and BD are on the opposite side of translocation domain (Kumaran et al. 2009)

chain and light chain of TeNT are nontoxic. But there is a second protease-sensitive site in heavy chain which splits the heavy chain approximately in half. In nonreducing condition, two chains can be isolated: one consists of 50 kDa heavy chain linked to 50 kDa light chain (referred to as fragment B), and other contains rest of heavy chain (50 kDa, fragment C). Fragment B is toxic to animal, and like botulinum toxin, it induces flaccid paralysis (Monteccuco and Schiavo 1994).



**Fig. 5** Structure of botulinum neurotoxin complex. The toxins are produced by bacteria as progenitor toxins, composed of toxin molecule associated with NBP and other NAPs (hemagglutinins or HAs). Based on the organization of these proteins, botulinum toxins are termed as LL (19S), L (16S), and M (12S) toxin. HA35 has the highest molar ratio in complex compared to toxin. According to Innoue et al. (1996), molar ratio of toxin to HA35 is 1.00:3.99 and 1.00:7.76 in 16S and 19S toxin, respectively

## 4.1 Neurotoxin-Associated Proteins

In bacterial cultures, BoNTs are released in the form of multimeric complexes, with the set of nontoxic neurotoxin-associated proteins (NAPs) coded by genes adjacent to neurotoxin gene. These complexes are termed progenitor toxins (Innoue et al. 1996; Minton 1995). Some BoNT-associated proteins have hemagglutinating activity (Fujita et al. 1995). A large nontoxic non-hemagglutinating protein of 139 kDa coded for the gene by a gene upstream to the BoNT gene is present in all seven serotypes of BoNTs called neurotoxin binding protein (NBP) (Innoue et al. 1996; Minton 1995; Fujita et al. 1995; Singh et al. 1995). One of the interesting facts about the NBP is that it has similar structural domains as BoNT (Gu et al. 2012). Three forms of progenitor toxins have been characterized: extra-large size (LL complex, sediments at 19 S, about 900 kDa), large size (L Complex, sediments at 16 S, about 500 kDa), and medium size (M complex, sediments at 12 S, about 300 kDa). The M complex is BoNT-NBP complex; L Complex of BoNT-NBP, and several components of NAPs with hemagglutinin activity, and the LL complex are believed to be a dimer of L complex associated with each other (Innoue et al. 1996). The four hemagglutinin (HA) proteins found in BoNT/A complex are HA15/17(15 or 17 kDa), HA33/35 (33 or 35 kDa), HA19/20 (19 or 20 kDa), and HA70 (70 kDa). Notably, HA52 and HA15/17 are fractions of HA70 (Fig. 5). Innoue et al. (1996) determined the molar composition of different protein in complex. HA33/35 has been purified from BoNT/A complex (Fu et al. 1998), and also has been expressed as a recombinant protein.

While BoNT/A exists in three complex forms, M, L, or LL, types B, C, and D BoNTs exist in two forms, L and M. Type E BoNT is now known to exist in L and

M forms, although NAPs composition of the L form is different (Singh and Zhang 2004; Kukreja and Singh 2007). Type F BoNTs are reported only in M form, and type G exists only in L form (Fujita et al. 1995, Fig. 5).

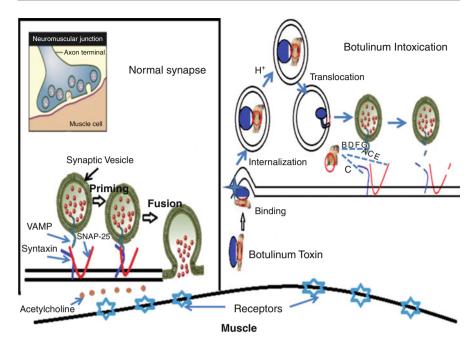
Since every BoNT serotype is associated with one or more NAPs, it is also important to examine biological role, if any, of these proteins. BoNTs in complex forms are more stable than in purified form. NAPs protect BoNTs against proteolysis and denaturation (Chen et al. 1998; Sakaguchi 1983; Kukreja and Singh 2007). Survival of the complex in harsh GI-tract conditions is believed to be a major reason for BoNTs to be food poison. Oral toxicity of BoNTs is more in complex form than their purified forms (Sakaguchi 1983). At least one of the NAPs has been shown to exhibit a series of biological functions. It has been shown to possess heat-shock protein characteristics (Shukla and Singh 1999, 2009) and is believed to act as molecular chaperone (Shukla et al. 1997). HA33 is also known to block neuronal apoptosis (Kumar et al. 2012). Furthermore, HA33 is known to assist in the translocation of BoNT across the gut wall after ingestion of the BoNT complex (Matsumura et al. 2008; Fujinaga et al. 2009). These results indicate that associated protein play an important role in the toxico-infection process of botulism.

# 5 Mechanism of Action of Clostridial Neurotoxin

It has been proposed (Simpson 1986, 2004; Hilger and Bigalke 1995) that the biological action of clostridial neurotoxins involves three steps: (a) binding and internalization of the neurotoxin via the HC through receptor-mediated endocytosis (cellular process by which cell absorbs and engulfing external molecule), (b) membrane translocation of LC into the cytosol purportedly through a pH-induced membrane channel formed by the HC, and (c) intracellular blocking of neurotransmitter release by proteolysis of one of the three SNARE proteins of by Zn-endopeptidase activity of the LC (Fig. 6). Isolated LC cleaves SNARE protein but it cannot readily internalize on its own and therefore remains nontoxic. Also, isolated  $H_C$  or  $H_N$  cannot cleave SNARE proteins, so remain nontoxic. So only when three domains are together they create the most toxic substance known to humanity. Interestingly, this unique assembly allows us to explore the possibility of combinatorial profiling of multidomain assemblies, for example, LC-TD for targeting LC function (Darios et al. 2010).

## 5.1 Receptor Binding and Internalization

As described above, BoNTs have trimodular architecture, and this unique architecture has a physiological meaning. This unique structural organization helps it to achieve its goal in harsh cellular environment of host. Its mechanism of action starts with the high-affinity interaction with a surface protein receptor, via ganglioside coreceptor. So far, protein receptors identified are SV2A for BoNT/A (Dong et al. 2006; Mahrhold et al. 2006); glycosylated SV2A and SV2B for BoNT/E



**Fig. 6** Mode of action of botulinum neurotoxin. *Left*, pre and postsynaptic events in normal neuronal cells. *Right*, steps involved in botulinum intoxication; binding, internalization, translocation

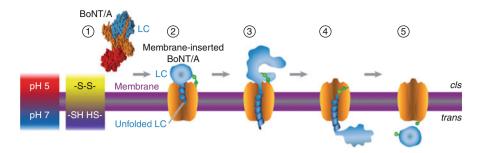
(Dong et al. 2008); glycosylated SV2A, SV2B, and SV2C for BoNT/F (Fu et al. 2009; Rummel et al. 2009); and synaptotagmin I and II for BoNT/B (Dong et al. 2003; Nishiki et al. 1994, 1996) and BoNT/G (Rummel et al. 2004a). Interestingly, the BoNT serotypes which exhibit highest sequence similarity in binding region share the same protein receptor, i.e., A, E, and F bind to SV2 and B and G bind to SytI and SytII. The protein receptor(s) for BoNT/C and BoNT/D remain unknown. Binding of BoNTs to the peripheral neuromuscular junction is the initial step of intoxication. Apart from these receptors BoNTs have high-affinity (with nM dissociation constants) binding with ganglioside co-receptor (Binz and Rummel 2009). BoNT types A, B, C, and F bind GT1b, GD1a, and GD1b, respectively. BoNT/E binds GT1b and GD1a, whereas BoNT/G recognizes all gangliosides with similar affinity. BoNT/D binds phosphatidylethanolamine (PE). The ganglioside binding pockets lie in the Hcc domain. Crystal structure of Hcc fold is highly conserved for all BoNTs (~60 % homology). GT1b binds on a crevice formed by Trp1266 and Tyr1267 on one face and Glu1203, His1253, and Phe1252 on the other face. Generally, lactose-binding motif has all these key residues (H...SXWY...G) (Rummel et al. 2004b), and this is conserved among all BoNT serotypes (Stenmark et al. 2008). This particular motif provides hydrophobic character to the binding cavity. In contrast, BoNT/D, which interacts with PE, (phosphatidylethanolamine) lacks the key residues and shows a dependency on Lys1117 and Lys1135 for PE

binding. Study of Hcc structure in the presence and absence of the SytII peptide (Jin et al. 2006; Rummel et al. 2004b) is similar. Also in the presence and absence of GT1b, no change in conformation was inferred (Stenmark et al. 2008), which implies not only the Hcc fold is rigid but that the binding sites for GT1b and SytII are nonoverlapping. This still needs to be looked at carefully as there is no structure available for a BoNT/A-SV2 complex or of a BoNT/B-GT1b complex. Nevertheless, we have BoNT/A Hcc-GT1b complex (PDB 2VU9) and BoNT/B Hcc-SytII peptide complex (PDB 2NM1), which allow us to understand the lipid and receptor binding. It is also believed that solvent accessible surface of each BoNT serotype is unique. Each ternary complex (BoNT/A-SV2-GT1b or BoNT/Eglycosylated SV2A-GT1b) may expose distinct charge groups. PIPs (phosphatidylinositol 4,5-bisphsophate) are minor phospholipids in cell membrane and involve in recruitment/regulation of protein in membrane and in signaling. PIP binding site is identified in BoNT/A H<sub>CN</sub>. Existence of two lipid binding sites, PIP and GT1b, suggests a dual anchorage mechanism for binding and internalization of this toxin (Muraro et al. 2009). Together with phospholipids, toxin acquires a new surface that confers a propensity for novel interaction which assisted in membrane insertion. Evidences suggest that after binding toxin does not enter the cell directly through plasma membrane, but is endocytosed into endosomes (enter into a lumen vesicular structure) in a temperature- and energy-dependent process (Dolly et al. 1984; Critchley et al. 1985; Black and Dolly 1986; Schiavo et al. 2000). The Hc domain of toxin plays an important role in internalization.

## 5.2 Pore Formation and Translocation

Second step after receptor-mediated endocytosis and internalization is the translocation of LC into the cytosol. LC must cross the hydrophobic barrier to reach cytosol. It is important to note that internalization and translocation both are distinct steps of cell intoxication. Translocation is a crucial step in the neurotoxicity of BoNTs. It is believed that  $H_N$  is translocation domain.  $H_N$  exhibits high-sequence conservation among all the serotypes with exception of the belt region that diverges significantly (Lacy and Stevens 1999).  $H_N$  helps in making pore in the endosomal membrane to facilitate LC translocation. LC translocation by HC stringently depends on the following conditions: (a) a pH gradient, acidic on the inside and neutral in the cytosol; (b) a redox gradient, oxidizing on the inside and reducing on the cytosolic side; and (c) a transmembrane potential (Montal 2010; Koriazova and Montal 2003; Fischer and Montal 2006, 2007).

There is no general consensus on how the translocation of LC occurs. According to one hypothesis, the LC unfolds at low pH and permeates through the transmembrane pore formed by  $H_N$ . After exposure to the neutral pH of the cytosol, the LC refolds, and it is released from the vesicle by reduction of the interchain disulfide bond. This is known as "tunnel" model (Boquet et al. 1976; Montecucco and Schiavo 1995), and formation of transmembrane ion-conducting pore is the pre-requisite for this model. This model has some problems: (1) the light chains of toxin



**Fig. 7** Schematic representation of BoNT/A LC translocation. BoNT/A prior to insertion (1), membrane inserted BoNT/(2), partial unfolding of light chain and translocation (3) and (4), and exit to cytosol (5) (Copyright (2007) O National Academy of Science, USA)

may be exposed to the fatty acid chains of lipids, and (2) channel activity values of the order of a few tens of picoseconds do not account for the dimensions expected for a protein channel that has to accommodate a polypeptide chain with lateral groups of different volume, charge, and hydrophilicity.

A second model, proposed by Beise et al. (1994) suggests a decrease in the internal pH of endocytotic vehicle carrying toxin by ATPase proton pump, followed by neurotoxin insertion into the lipid bilayer. Electrochemical gradients and osmotic lysis cause destabilization of the lipid bilayers, breaking the membrane barrier, and the cargo of toxin molecules are released into the cytosol. This model explains the translocation of toxins fairly well, but some experimental findings are inconsistent with it. It is already shown that TeNT does not lyse the plasmalemma of neuronal cell at pH 5.0 and also diphtheria toxin forms channel without lysing the cell (Beaumelle et al. 1992).

Third model is the cleft model (Humeau et al. 2000; Simpson 2004) in which there is conformational change at low pH in both HC and LC in such a way that they expose their hydrophobic surfaces and enter into lipid bilayer by getting in contact with its hydrophobic core. The HC forms a transmembrane hydrophilic cleft that nests the passage of the partially unfolded LC with its hydrophobic segments facing the lipids. This process may be driven by electrochemical gradient, such as pH gradient across endosomal membrane. Neutral pH of cytosol and/or cytosolic chaperones help LC to fold and regain its water-soluble neutral conformation after reduction of the interchain disulfide bond. In the cleft model, the ion channel is a consequence of membrane translocation rather than a prerequisite.

A recent study by Montal (2010) suggests a very interesting mechanism of translocation, suggesting the role of interchain disulfide bond. This explanation suggests occlusion of heavy chain channel by light chain, if disulfide bond is reduced prior to translocation. Also, disulfide bond reduction within the bilayers aborts translocation. Translocation is only complete when disulfide bridge and the C-terminus of the LC enter the cytosol. According to this observation, it may be suggested that channel formation and translocation are two different events and not associated with each other (Montal 2010, Fig. 7).

Examination of BoNT crystal structure also reveals a transmembrane region (548–872) and a loop (561–568; completely disordered) connecting translocation belt (492–545 in BoNT/A and 481–532 in BoNT/B) and HN, which may change its conformation at low pH. The translocation domain appeared to be sensitive to environment variables, as the pH of this region is 4.66 (as compared to 9.1 for the binding domain and 6.0 for the whole toxin). There are several clusters of negative charges as well as proximal flexible structures in HN which can assist the pore formation (Lacy and Stevens 1999).

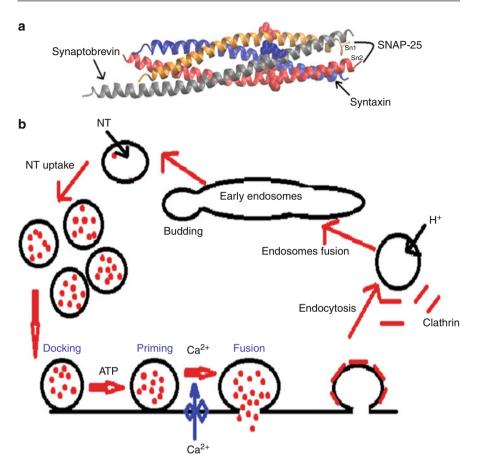
Evidences show that there are similarities in the action of botulinum and tetanus neurotoxins. TeNT, like BoNT, is a 150 kDa single-polypeptide protein with a heavy chain (100 kDa) and a light chain (50 kDa) region held together by a disulfide bridge and by noncovalent protein-protein interactions; TeNT also binds to peripheral nerve terminals, internalizes through endocytosis, and retrogrades to central nerves, where the light chain translocates into cytosol, leading to biochemical action that blocks neurotransmitter release.

The carboxyl terminal of TeNT heavy chain binds to cholinergic receptors of the presynaptic membranes of the central and peripheral nervous system, and possibly to sympathetic adrenergic fibers. Polysialogangliosides are an example of cholinergic receptors. TeNT binds tightly to such receptors because of the protein's multiple interactions with sugar and protein binding sites. It is retrogradely transported within the axon to the cell body. TeNT is released from the postsynaptic dendrites and taken up by the presynaptic nerve terminals. The entire journey of TeNT from peripheral neurons to its ultimate site of action, inhibitory neurons of central nervous system, involves binding, internalization, and transport within a cell. Transport of the toxin seems to occur within smooth vesicles, cisternae, and tubules. The toxin acts specifically on the interneurons of the spinal cord that inhibit voluntary muscle action. The N-terminus of the heavy chain is the translocation domain because it forms pores or toxin channels in the vesicle lipid bilayers of spinal cord neurons. The light (50 kDa) chain of TeNT is the toxic part of the protein. This portion is a specific protease that cleaves synaptobrevin, a vesicleassociated membrane protein (VAMP), at a single site. VAMPs help mediate membrane fusion, vesicle transport, and neurotransmitter release. Cleavage of synaptobrevin would not only denature the protein, but it would also prevent the exocytosis of the glycine and  $\gamma$ -aminobutyric acid neurotransmitters that control muscle contraction. With such actions inhibited, the nerve cells that control voluntary muscle action fire constantly and lead to the characteristic unopposed muscle spasm leading to spastic paralysis.

# 5.3 Blockage of Neurotransmitter Release

#### 5.3.1 SNAREs and Neurotransmission

To understand the intracellular mechanism of action of clostridium neurotoxins, we first need to understand its substrate SNARE proteins and its role in neurotransmitter release. SNAREs are set of three proteins: synaptic vesicle protein, VAMP



**Fig. 8** (a) Tetrahelical assembly structure of SNARE complex, (b) schematic representation of the processes of endocytosis and exocytosis. Docking, priming, and fusion of exocytosis are represented in this figure. SNARE proteins (syntaxin, SNAP-25, and synaptobrevin) are critically involved in different steps of the exocytosis process. Synaptic vesicles loaded with neurotransmitter dock near the plasma membrane. Partial SNARE complex formation occurs in the priming step. Primed vesicles fuse very rapidly with the plasma membrane after calcium influx upon the arrival of the action potential. Energy for the fusion comes from SNARE assembly

(vesicle-associated membrane protein also called as synaptobrevin), and plasma membrane proteins, syntaxin, and SNAP-25 (synaptosome-associated protein of 25 kDa) (Fig. 8a). These three proteins are membrane receptors for NSF (*N*-ethylamide-sensitive factors) and SNAPs (soluble NSF attachment proteins and are named as SNARE (SNAP receptors) proteins. SNAREs can be divided into two categories based on their localization inside the cells: v-SNAREs, localized in vesicle membrane, and t-SNAREs, localized in target membrane or plasma membrane. They are also sometimes termed as R-SNARE (middle group arginine) and Q-SNARE (middle group glutamines). SNARE proteins are characterized by

a conserved 60-70 amino acid motif. In addition to this motif SNAREs also have sequences that anchor them to the membrane. Syntaxin and synaptobrevin are anchored through their respective transmembrane domains, whereas SNAP-25 associates with the plasma membrane via the palmitovlation of cysteine residues in the linker region. These proteins play a very important role in the membrane fusion and neuronal exocytosis, one of the important physiological processes in neuronal cells. SNAP-25 and syntaxin are highly conserved in their isoforms. SNAP-25 is also needed for axonal growth and regeneration (Osen et al. 1993). One of the unique characteristic features of SNARE proteins is their assembly in 1:1:1 stoichiometry to form a very stable ternary complex, which is resistant to dissociation by sodium dodecyl sulfate (SDS). During the ternary complex formation, carboxy-terminal domain of syntaxin (amino acids 199–243; H-3 helix) binds to SNAP-25. Complex of SNAP-25 and syntaxin binds to the central part of the synaptobrevin (amino acids 27–96). All of SNAP-25 is required for binding with synaptobrevin (VAMP2, 29-60) and amino- and carboxy-terminal domains participate in binding to syntaxin (syntaxin 1, 199–243). Assembly of SNARE complex brings the two opposing membrane bilayers of synaptic vesicle and plasma membrane closure and acts as a bridge. The arrival of action potential triggers inflow of Ca<sup>2+</sup> ion, allowing the vesicle to dock at the active zone. After docking, second Ca<sup>2+</sup> influx in the active zone creates a fusion pore involving two other proteins: synaptophysin and physophillin. Other vital components in this process are synaptotagmin and complexin. Complexin interacts with tetrahelical strands (Fig. 8a) of SNARE complex and prevents neurotransmitter release until an action potential arrives at synapse to facilitate quantal release (Kummel et al. 2012). Synaptotagmin binds to Ca<sup>2+</sup> ion and acts as a sensor for vesicle fusion and quantal neurotransmitter release (Kummel et al. 2012). Formation of initial pore leads to complete fusion of vesicle membrane to plasma membrane, and the content of vesicle (neurotransmitter) is released into the synaptic cleft. The whole process is called exocytosis (Fig. 8a and b).

## 5.3.2 Motifs and Conformations Involved in Unique Substrate Recognition of BoNT Endopeptidase

All the light chains are Zn-metalloprotease of thermolysin family but are unique among Zn proteases due to their stringent specificity for substrates or at least long peptide segments containing highly selective cleavage sites. BoNT LCs require 16 to >50 residue long substrates whereas other Zn proteases can hydrolyze as short as two residue peptides (Shone et al. 1993; Cornille et al. 1997; Shone and Roberts 1994). The known substrate for BoNTs is SNARE proteins with stringent substrate specificity. TeNT and BoNTs target and peptide bond specificity is displayed in Table 2.

Each of the clostridial toxins cleaves their targets at distinct sites, except BoNT/ B and TeNT which cleave the same peptide bond on the same substrate. This exquisite selectivity is believed to be due to the presence of highly conserved motifs in SNARE proteins. These motifs are termed as S1 to S4 in SNAP-25; V1 and V2 in VAMP; and X1 and X2 in syntaxin. Although BoNT/B and TeNT attack the same

Table 2     TeNT and BoNT			
	Toxin type	Substrate	P4-P3-P2-P1-P1'-P2'-P3'-P4'
target and peptide bond specificities. Cleavage site is	TeNT	VAMP	G- A- S- Q- F- E- T- S
at P1–P1'	BoNT/A	SNAP-25	E- A- N- Q- R- A- T- K
	BoNT/B	VAMP	G- A- S- Q- F- G- T- S
	BoNT/C	SNAP-25	A- N- Q-R- A- T- K-M
	BoNT/C	Syntaxin	D- T- K- K- A- V- K- F
	BoNT/D	VAMP	R- D- Q- K-L- S- E- L
	BoNT/E	SNAP-25	Q- I- D- R- I- M- E-K
	BoNT/F	VAMP	E- R- D-Q- K-L- S- E
	BoNT/G	VAMP	E- T- S- A- A-K- L-K

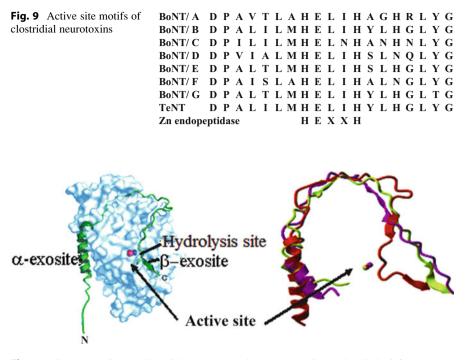
Gln-Phe peptide bond, these toxins display distinct affinities for V1 and V2. For efficient cleavage, BoNT/A and BoNT/E require C-terminal region of SNAP-25 which has S4 motifs. However, S1 (leu20-Arg30), S2 (leu36-Ile45), and S3 (Met50-Glu59) motives partially substitute for S4 (Glu145-Ser154) motif (Washbourne et al. 1997). SNARE proteins are protected from proteolytic attack if they are bound to ternary complex (Poulain et al. 2008). This is probably due to masking of the cleavage site or recognition motifs.

Several evidences suggest that neurotoxins recognize tertiary structure of their substrates:

- (a) Clostridial neurotoxins require long substrates compared to other Zn-metalloprotease.
- (b) They cleave their substrates exclusively at single-specific site out of several identical peptide bonds present in their respective target proteins.
- (c) Synthetic peptides of same cleavage segments are not hydrolyzed by the neurotoxins; they are shown to be good inhibitors of proteolytic activity of the toxins. This means although they can bind, they are not cleaved by toxins (Schiavo et al. 1992a, b, 1993a, b; Dayanithi et al. 1994).
- (d) Light chains of these toxins cleave very specific bonds even if the same bonds are present in other part of the substrate sequence (Gln15-Arg16 of SNAP-25 in case of BoNT/A and Arg59-Ile60 and Arg191-Ile192 in case of BoNT/E).

Apart from SNARE, several homologues of these proteins are known as a substrate for BoNT, which can be cleaved to different degrees. Murine SNAP-23 can be cleaved by BoNT/E and to some extent by BoNT/A, but human SNAP-23 is resistant to both because of mutation Lys185Asp and Pro185Arg (Washbourne et al. 1997; Vaidyanathan et al. 1999; Pickett and Perrow 2011). Mutations in amino acid sequences of the toxin targets make them resistant to proteolysis and can account for almost all cases of species insensitivity to BoNT or TeNT.

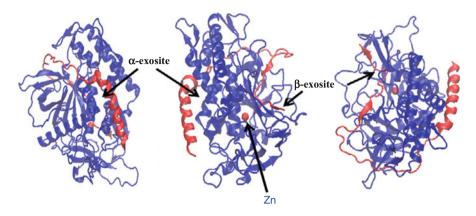
LC of BoNT is a compact globular structure composed of a combination of  $\beta$ -sheets and  $\alpha$ -helices (PDB ID 1XTG; 1 F82; 1T3A; 2ISE). All the crystal structures reported so far are derived from a truncated version of the LCs.



**Fig. 10** Structure of BoNT/A LC in complex with truncated SNAP-25 (Sn2, *left*), showing interaction of SNAP-25 (Sn2 segment) with  $\alpha$ - and  $\beta$ -exosites and active site. *Right* figure shows overlapped structure of SNAP-25 (Sn2 segment, *red*) and HC belt of BoNT/A (*magenta*) and BoNT B (*lime*) (Adapted from Brunger et al. 2007)

The central part of this enzyme is  $Zn^{2+}$  binding motif, HEXXH + E, and is conserved among all BoNTs. The finding is that all the clostridial neurotoxins have similar active site motifs, suggesting common evolutionary origin of these proteins (Fig. 9).

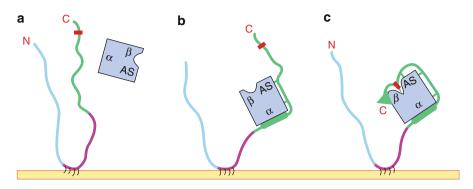
It is notable that the active site of BoNT is located in a very deep and narrow crevice, whereas in thermolysin active site is broad and very near to the surface (Fig. 10). In case of BoNT/A LC, active site crevice travels through half of the enzyme structure (20 Å deep) and is negatively charged (Brunger et al. 2007). Interestingly, translocation domain has a belt, a unique loop in extended conformation (residues 492–545 for BoNT/A and 481–532 for BoNT/B). This belt has a remarkable similarity with residues 141–205 (the Sn2 segment) of SNAP-25 in spite of low sequence similarity of the belts among all the clostridial toxins (BoNT and TeNT; Lacy and Stevens 1999; Chen et al. 2007b, Fig. 10). For BoNT/A, the catalytic Zn<sup>2+</sup> is bound to the active site by interaction with imidazole ring of His223 and His227 and with the carboxyl side chain of Glu262. The Zn<sup>2+</sup> is also coordinated by a water molecule, which is activated by Glu224, and believed to



**Fig. 11** Crystal structure of BoNT/A LC with its substrate SNAP-25 (Sn2).  $\alpha$ -exosite and  $\beta$ -exosite are shown as a primary substrate recognition point. 250 and 370 loops are also interacting with substrate (Breidenbach and Brunger, Nature 2004)

play a critical role in proteolysis. However, Glu262, which is directly coordinating to active site Zn, also plays a pivotal role in the stability of active site (Kukreja et al. 2007). Furthermore, four flexible loops, 50/60 loop, 170 loop, 370 loop, and 250 loop, form the rim of active site cleft and are believed to take part in substrate binding. Rest of the protein is a mixture of  $\alpha$ -helix and  $\beta$ -sheets.

A topographical model for the interaction of BoNT with its substrate has been described by Breidenbach and Brunger (2004, Fig. 11) who used enzymatically inactive version of BoNT/A LC (created by two point mutations, E224Q and Y366F) and carboxy-terminal SNARE domain of SNAP-25, residues 141-204 (Sn2), to obtain crystal structure of enzyme-substrate complex. The crystal structure of the complex revealed that apart from active site, this molecule has two exosites:  $\alpha$ -exosite and  $\beta$ -exosite.  $\alpha$ -exosite is a hydrophobic patch formed at the interface of four light chain- $\alpha$ -helices, residues 102–113 ( $\alpha$ -helix 1), 310–321 ( $\alpha$ -helix 2), 335–348 ( $\alpha$ -helix 3), and 351–358 ( $\alpha$ -helix 4).  $\beta$ -exosite is a distorted two strand in the 250 loop of the LC, a peptide segment with residues 242–249. The amino-terminal helical region of Sn2 interacts with the endopeptidase  $\alpha$ -exosite, whereas C-terminal of Sn2 interacts with β-exosite. The hydrophilic face of the amphipathic Sn2 helix is oriented towards the solvent, while the hydrophobic side chain buried against the  $\alpha$ -exosite. Interaction with  $\beta$ -exosite is primarily mediated by backbone interactions, but Met 202 of Sn2 is also very critical for this interaction. Substrate shorter than residues prior to Met 202 cannot be cleaved by BoNT at a measurable rate (Schmidt and Bostian 1995). β-exosite is very critical for the recognition of BoNT. Notably, residues 183–190 of Sn2 detach from the surface of LC and rejoin the LC at Arg191. There are series of anchoring points that could serve as additional determinants of substrate specificity. Substrate residues 170-172 and 192-193 are two anchoring points, and individual side chains from Arg176, Ile178, Ile181, and Glu183 are also recognized by the endopeptidase. This may be one of the reasons for the requirement of long substrate. The largest



**Fig. 12** Schematic model of BoNT/A LC substrate recognition and cleavage (Adapted from Breidenbach and Brunger 2004). (**a**) Sn1 (*cyan*) and Sn2 (*green*) domains of SNAP-25 are attached to presynaptic membrane by palmitoylation (*black*) of cysteine residues of SNAP-25. (**b**) Binding of BoNT/A is initiated by recognition of Sn2 domain of SNAP-25 by  $\alpha$ -exosite of protein and few anchoring points (*green* notches). (**c**) After these interaction SNAP-25 binds to  $\beta$ -exosite, inducing conformation change in 370 loop, which leads to cleavage of SNAP-25 by BoNT/A LC

structural changes that are related to substrate binding occur in 250 loop ( $\beta$ -exosite) and 370 loop In BoNT/A; the 250 loop adopts an open conformation which contracts and folds over the 370 loop upon substrate binding at the  $\beta$ -exosite.

The model of active site and exosites of BoNT/A endopeptidase explains the mechanism of SNAP-25 substrate cleavage as follows. SNAP-25 is attached to presynaptic membrane through palmitoylation of cysteine residues. C-terminus and N-terminus are unstructured or flexible in the uncomplexed SNAP-25. SNAP-25 upon binding to BoNT/A probably initiates helix formation which is recognized by  $\alpha$ -exosite and anchored points and finally by  $\beta$ -exosite (Figs. 11 and 12).

A major point against this model is that the BoNT/A L chain and SNAP-25 co-crystal structure do not show any significant direct interaction of the  $\alpha$ -exosite of the light chain with the S4 motif of SNARE, not considered critical for binding of SNAP-25 (Breidenbach and Brunger 2004), but are critical for the endopeptidase activity (Washbourne et al. 1997). Thus, while the crystal structures provide useful information on the folding patterns of BoNTs, and on the interaction between BoNT/A light chain and SNAP-25, these do not take into consideration the role of dynamic structures in the functioning of the molecule. Nevertheless, this cocrystal structure provides a very good model to understand the behavior of this molecule with its substrate. The crystal structure explains why clostridium neurotoxins can cleave free SNAREs but not the SNARE complex because when SNARE complex is formed, then the residues of SNAP-25 involved in recognition/bindings are not accessible for these interactions. Notably, C-terminus of BoNT/A LC plays important role in catalysis. Finding by Baldwin et al. (2004) suggests that minimum residue needed for optimal catalysis is 1–425. Deletion of C-terminal region beyond 425 has a small effect on substrate binding but drastically reduced the catalytic efficiency. It has been speculated that there is extensive H-bond network in the

C-terminal region which stabilizes the structure of C-terminal loop region as well as active site core (Baldwin et al. 2004).

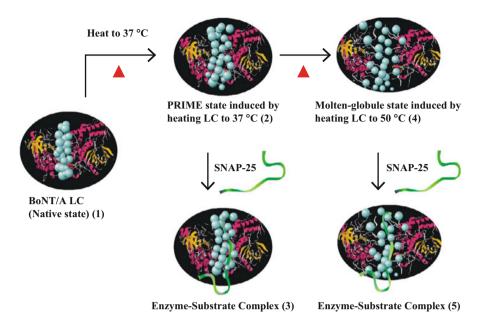
## 5.3.3 Static Versus Dynamic Structure: Role of PRIME Conformation

The static crystal structure of botulinum endopeptidase determined by x-ray diffraction studies (Segelke et al. 2004; Breidenbach and Brunger 2004) does not adequately describe the dynamic properties of the enzyme in solution where it represents a manifold of conformational substates. Indeed, crystal packing forces might lead to structural arrangements, which might not be relevant to solution conditions. Conformational fluctuations resulting from collective modes of motion, which involve concerted displacements of many atoms, can push the unbound states of enzymes into conformations closely resembling the bound states, thereby priming them to form complexes with the substrate. Hence, every conformational state of the enzyme that is populated along the reaction pathway presents a unique opportunity for interactions with substrate or drug molecules.

We have recently discovered biologically active molten globule structure of the BoNT/A (Cai and Singh 2001). The catalytic domain of BoNT/A also possesses a novel conformation at the physiological temperature of 37 °C and is named as PRIME (pre-imminent molten globule enzyme) state (Kukreja and Singh 2005). The PRIME state is dynamically flexible and is likely to facilitate specific interactions with its substrate, SNAP-25 for its optimum and selective enzymatic activity. Thus, we suspect that the PRIME conformational state provides a flexible structure which plays a very critical role in the biological function of BoNT/A, especially in its intracellular toxic action.

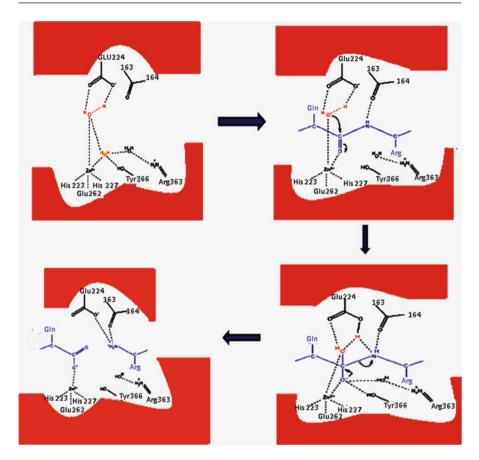
The schematic model of the temperature-induced PRIME conformational and molten globule states of BoNT/A light chain, as represented in Fig. 13, reflects how these states in BoNT/A light chain may facilitate binding and cleavage of SNAP-25. The intramolecular mobility of PRIME state is significantly higher than that of the native state which can be attributed to increased dynamics and expansion of the protein core, facilitating a maximum specific binding of SNAP-25, leading to its cleavage. The molten globule conformation observed at 50 °C, on the other hand, has a considerably intact secondary structure, loose packing of side chains in the protein core, and partial unfolding of loops. This structure also binds to SNAP-25, although to a lesser degree than the PRIME state, thus showing only about 61 % of the optimum enzymatic activity (Fig. 13).

Our observation of the PRIME conformation of BoNT/A LC with optimum enzymatic activity is the first observation of enzymatically active molten globule or molten globule-like structure (Kukreja and Singh 2005). The role of such a dynamic structure is more significant in view of the extreme specificity of the endopeptidase towards its substrate, SNAP-25, and its tremendous utility in designing specific antidotes against botulism threats. This structure of BoNT/A light chain seems to play a major role in its endopeptidase activity and particularly as it interacts with potential inhibitor candidates. Currently, BoNT/A light chain used in most



**Fig. 13** Schematic model representing the PRIME state and the molten globule state in BoNT/A LC facilitating binding of substrate SNAP-25. The *blue*-colored balls represent the nonpolar side chains which are tightly packed in the native conformation (1). Upon heating to 37 °C, there are significant alterations in the polypeptide folding and the protein core becomes slightly loosened compared to the native state and forms the PRIME state (2), facilitating binding of SNAP-25 to form the enzyme-substrate complex (3). BoNT/A LC exhibits optimum activity in the PRIME state. Further heating of BoNT/A LC to 50 °C leads to the formation of the molten globule (4), facilitating binding of SNAP-25 to form the enzyme-substrate complex (5) although to a lesser degree than the PRIME state, thus exhibiting only about 61 % of the optimum enzyme activity (Kukreja and Singh 2005)

laboratories working on structure and inhibitor development is a truncated form of the 448 residue full-length polypeptide. The forms being used are LC-444, LC-438, LC-425, and LC-424. Although all these truncated forms have enzyme activity, their interactions with potential inhibitors as well as their substrates (peptide substrate and whole protein substrate, SNAP-25) were peculiarly different (Baldwin et al. 2004; Fredenburg et al. 2007), and a dynamic conformational variability could explain these differences. Silvaggi et al. (2008) have in fact observed limited flexibility of active site, although those changes would not necessarily explain differential interaction of BoNT endopeptidases with their respective substrates, because of the involvement of exosites in substrate binding (Breidenbach and Brunger 2004). Burnett et al. (2007) have observed that "Our inability to identify an acceptable binding mode for mpp-RATKML led to a pivotal question: is it possible that the conformation of the x-ray determined BoNT/A LC may not be the bioactive conformation, at least with respect to inhibitor binding?"



**Fig. 14** Schematic representation of proposed BoNT/A LC mechanism. A polarized water molecule, by Glu223 and  $Zn^{2+}$  (*top left*), nucleophilically attacks the carbonyl carbon of the scissile bond to form an oxyanion (*top right*). This oxyanion is stabilized by Tyr366 (*bottom right*). Peptide bond of the substrate is cleaved by two proton transfer for activated water molecule mediated by Glu223. This process resulted in a protonated amine (*bottom left*)

Mechanistic details of a flexible structure interacting with its whole substrate have yet to be worked out, but when that is resolved it could reveal transformative role of protein dynamics in the protein-protein interactions involved in such a high specificity in BoNT/A endopeptidase-SNAP-25 interactions.

## 5.3.4 Chemical Mechanism of Substrate Cleavage

Proposed mechanism of substrate cleavage is depicted in Fig. 14. In the first step water molecule is polarized by Glu224 and  $Zn^{2+}$ . Activated oxygen of water mounts a nucleophilic attack on the carbonyl carbon of P1 of scissile bond. Side chain of Tyr366 is responsible for stabilization of the resulting oxyanion, which develops at

P1 carbonyl oxygen atoms. Oxyanion is also stabilized by Arg363 through a water molecule. Peptide bond cleavage is likely achieved by two proton transfer from the attacking water molecule mediated by Glu224 and results in the protonated amine of P1'. Proposed mechanism is slightly different from the mechanism of another Zn-metalloprotease, thermolysin. In thermolysin, Tyr157 and His231 have been proposed to stabilize the transition state (Matthews 1988). No residues have been identified to involve in the stabilization of the transition state of ES (enzyme-substrate) complex of BoNT-catalyzed proteolysis. In fact, the space occupied by these two residues in thermolysin is occupied by Sn2 residues in the case of BoNT/A. This leads to two hypothesis: either there are no residues in the case of BoNT/A to stabilize the evolving oxyanion or that BoNT/A might involve substrate-assisted catalysis.

His223, Glu224, His227, and Glu262 are the important residues in BoNT/A active site, and they are conserved in the other serotypes. Mutation of Glu224 to Asp leads to almost total loss of activity, and removal of His227 abolishes the catalytic activity of BoNT/A. Analysis of TeNT neurotoxin has confirmed that these key residues are identical to BoNT/A. Other residues, such as Glu271 and Tyr375 (in TeNT) and Arg 363 and Tyr365 (in BoNT/A), have been shown to be involved in catalysis, and it has been proposed that these residues may be involved in transition state stabilization.

# 6 Effects and Consequences of Botulinum Neurotoxin Exposure

As mentioned earlier, botulinum toxins are the most poisonous substance known. The LD50, dose of these toxins in mouse when administered intravenously, subcutaneously, or intraperitoneally, is estimated 1 ng/kg. Tetanus toxin LD50 is 2 ng/kg. All the botulinum toxins are less toxic when exposure is through pulmonary route. Human LD50 is 3 ng/kg (McNally et al. 1994). The effect of botulinum toxin may affect areas away from the injection site and cause serious symptoms including loss of strength and all-over muscle weakness, double vision, blurred vision and drooping eyelids, hoarseness or change or loss of voice (dysphonia), trouble saying words clearly (dysarthria), loss of bladder control, trouble breathing, and trouble swallowing. While all the seven serotypes have similar domain organization and function, and show only immunological distinction (Simpson et al. 2004), duration of their intracellular action (longevity) and potency varies substantially. BoNT/A, BoNT/B, and BoNT/E intoxication is persistent for about 180 days, 90 days, and 30 days, respectively (Adler et al. 2001; Dolly and Aoki 2006; Pickett and Perrow 2011).

Persistence of BoNT/A intoxication is because of stability of its catalytic domain, light chain, inside the neuronal cells. This is supported by intoxication of BoNT/A in spinal cord neurons (Keller et al. 1999) and muscle paralysis experiments (Adler et al. 1996). Notably, neurons and neuroendocrine cells have high level of tyrosine kinases (Brugge et al. 1985; Maher 1988). Therefore, it has been

speculated that intercellular phosphorylation of light chain may contribute to latter's longevity (Montiel et al. 1996). Recently, it has also been proposed that differential ubiquitination of different BoNT light chains may be responsible for differential intercellular stability of BoNT proteins (Tsai et al. 2010). Because of the longevity of BoNT intoxication, and that too due to the survival of toxin inside the cell, an inadvertent consequence is the possibility of the host cellular response to the long-term presence of the toxin.

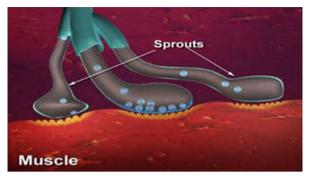
Microarray analysis reveals that 169 genes were upregulated, while 60 genes were downregulated in epithelial cell (HT-29), whereas 223 genes were upregulated and 18 genes were downregulated in neuronal cell model (SH-SY5Y) exposed to toxin for just 96 h. These modulated genes are associated with neuroinflammatory, ubiquitin-protease degradation, phosphatidylinositol, and calcium signaling in SH-SY5Y pathways (Thirunavukkarasu et al. 2011). Interestingly, these modulated genes are different from those observed upon exposure of HT-29 cells, which are model for gut epithelial cells. Are there gene expression responses related to survival of BoNT intracellularly? Further research work of cells to different serotypes of BoNT will shed more light on the genomic host response and BoNT longevity inside the neuronal cells.

As BoNTs are associated with NAPs, and in most cases of botulism cases or even BoNT as a therapeutic drug, patients are exposed to the BoNT complex, it is important to consider the effect of NAPs as well on biochemical and physiological host response. Immunogenicity of BoNT/A complex is more than that of pure BoNT/A itself. Interestingly, NAPs are more immunogenic than the toxin (Kukreja et al. 2009). HA33, molecule major component of BoNT/A NAPs, exhibits maximum immunogenicity. HA33 is also shown to inhibit apoptosis (Kumar et al. 2012). Apoptosis is also inhibited by BoNT/A in SH-SY5Y cells (Kumar et al. 2012), although BoNT/C is shown to enhance apoptosis in cerebral granular cells (Berliocchi et al. 2005). While these studies indicate that BoNT and NAPs significantly affect cell metabolism and physiology, there are many questions, including variations in responses to different BoNT types, variation in responses of different cell and tissue types, and variation in responses due to different states of cellular physiology, that remain to be addressed with further experiments.

Effect of botulinum toxin on nerve impulse activity and muscle contractions resumes over time as a result of sprouting near to nerve ending, thus renewing the ability of the nerve to cause muscle contraction. Eventually, the new nerve retracts and original nerve ending regains its function (Jankowic 2004; De Paiva et al. 1999, Fig. 15).

Release of neurotransmitter is affected by botulinum neurotoxins due to the cleavage of SNARE proteins, but that is not the only process affected by BoNT. Neuropeptide P, or substance P, involved in neurogenic inflammation and pain disorders is also affected by BoNT (Aoki 2004). There is lack of evidence of direct effect of BoNT on central nervous system. Because of its size, 150 kDa, BoNT cannot penetrate blood–brain barrier (Dressler et al. 2005). Possibility of retrograde transport has been suggested previously; however, no transsynaptic transport has





been observed. In vitro studies suggest that BoNT can affect central nervous system (Antonucci et al. 2009; Caleo and Schiavo 2009). BoNT/E has been shown to affect glutamate release and block spikes activity of pyramidal neurons in vivo (Bozzi et al. 2006). A recent study finds that BoNT spreads in CNS tissues (Jamie 2008); however, the injected dose was way too high compared to the doses used in therapeutic application.

# 7 Medical Applications of Clostridial Neurotoxins

Despite being highly toxic, clostridial neurotoxins are highly effective therapeutic agents. BoNT-based therapeutic products are approved by USFDA which include Botox<sup>®</sup>, Dysport<sup>®</sup> (BoNT/A complex), Myobloc<sup>®</sup>, Neurobloc<sup>®</sup> (BoNT/B complex), and Xeomin<sup>®</sup> (purified BoNT/A). These products are very popular cosmetic drugs in the world. USFDA also approved BoNT/A for the treatment of strabismus, blepharospasm, and hemifacial spasm. Since BoNT affects autonomous nervous system, it has been shown for effective treatments in hyperhidrosis, migraine, and myofascial pain. However, like other medical drug, BoNT treatment has some side effects too. Since these therapeutic agents are proteins, they have potential for antibody formation leading to a decrease in effectiveness of treatment. Since cholinergic neurons are important part of cognitive systems, BoNT can have potential role there as well and can have long-term effect on cognition (Lackovic et al. 2009; Ney and Joseph 2007). But this has been observed in relatively low number of patients.

# 8 Potential Future Applications

Each domain of clostridial toxins has very unique characteristics which can be used for some other medical applications. Retargeting of the endopeptidase domain of clostridial neurotoxin neuronal and nonneuronal cells is successfully performed. Retargeted  $LH_N$  (light chain fused with N-terminus of heavy chain) with nerve growth factor (NGF) and wheat germ agglutinin (WGA) has already been reported (Chaddock et al. 2000a, b). BoNT usefulness in alleviating pain is still under investigation. Mechanism is not very clear and there is no clear dose-response relationship. Retargeting technology has been exemplified using a variety of targeting ligands in a diverse range of cell types (Yeh et al. 2011; Pickett 2010; Somm et al. 2012; e.g., HEK (human embryonic kidney) 293 cells and BMDMs (bone marrow-derived macrophages). Additionally, the ability of BoNT and TeNT to transport large polypeptide makes them a good delivery vehicle. TeNT has been shown to transport DNA and enzymes in neurons (Toivonen et al. 2010; Wang et al. 2012). BoNT/A and BoNT/B have been tested for treatment of Parkinson's, rabies. and Tuorette's disease. One thing for sure is that anticholinergic effect of BoNT can be exploited for medicinal applications associated with physiological system employing cholinergic neurons. Above observations suggest that clostridial neurotoxins have become a valuable research and medicinal tool. Also it seems more uses of BoNT and its derivatives are in the offing. In conclusion, Mother Nature has provided a unique molecule with unique characteristics which can be exploited for other beneficial uses.

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# **Botulinum Neurotoxins as Therapeutics**

## Raja Mehanna and Joseph Jankovic

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#### Abstract

Since first recognized as the cause of food-borne botulism in the early nineteenth century, botulinum toxin was suggested as a potential treatment for involuntary spasms and movements. Multiple double-blind, placebo-controlled, and open-label studies provided evidence that botulinum toxin is a powerful therapeutic tool in a variety of neurologic and other disorders including ophthalmologic, gastrointestinal, urologic, orthopedic, dermatologic, secretory, painful, and cosmetic disorders. We here review the basic mechanisms of botulinum toxin action at the neuromuscular junction and discuss some of its main clinical applications.

R. Mehanna (🖂) • J. Jankovic (🖂)

Parkinson's Disease Center and Movement Disorders Clinic, Department of Neurology, Baylor College of Medicine, Houston, TX, USA

e-mail: doc\_raja\_mehanna@yahoo.com; josephj@bcm.edu

#### Keywords

Botulinum toxin • Clinical application • Mechanism of action

List of Abbrev	List of Abbreviations			
AAN	American Academy of Neurology			
Ach	Acetylcholine			
BoNT	Botulinum toxin			
CD	Cervical dystonia			
CD-PROBE	Cervical Dystonia Patient Registry for the Observation of			
	Botulinum Toxin Type A Efficacy Study			
EMG	Electromyography			
FDA	Food and Drug Administration			
nAChRs	Acetylcholine receptors			
OMB	Oromandibular dystonia			
PD	Parkinson's disease			
PREEMPT	Phase III Research Evaluating Migraine Prophylaxis Therapy			
SNARE	Soluble N-ethylmaleimide sensitive factor attachment protein			
	receptor			
TTA	Therapeutics and Technology Assessment			
UBI	Unilateral brow injection			
VAMP	Vesicle-associated membrane protein			

#### 1 Introduction

Since first recognized in 1817 by Christian Andreas Justinus Kerner as the cause of food-borne botulism, botulinum toxin (BoNT) was suggested as a potential treatment for involuntary spasms and movements. However, it was not until 1973 when Alan Scott, M.D., pediatric ophthalmologist and researcher in San Francisco, demonstrated that BoNT injections into the extraocular muscles improved strabismus in monkeys. The first report of clinical application of BoNT was published in 1984, when it was demonstrated to be safe and effective in the treatment of blepharospasm (Frueh et al. 1984). Subsequent multiple double-blind, placebo-controlled, and open-label studies provided evidence that BoNT was a powerful therapeutic tool in a variety of neurologic and other disorders including ophthalmologic, gastrointestinal, urologic, orthopedic, dermatologic, secretory, painful, and cosmetic disorders (Truong and Jost 2006; Jankovic 2009b; Hallett et al. 2013; Naumann et al. 2013) (Table 1).

We will first review the basic mechanisms of BoNT action at the neuromuscular junction and then discuss some of its main clinical applications.

Table 1 Clinical	Neurologic disorders		
applications of botulinum	Blepharospasm (lid "apraxia")		
toxin	Oromandibular–facial–lingual dystonia		
	Laryngeal dystonia (spasmodic dysphonia)		
	Cervical dystonia (torticollis)		
	Limb dystonia		
	Task-specific dystonia (e.g., writer's cramp)		
	Other axial/focal/segmental dystonias (primary, secondary)		
	Hemifacial spasm		
	Limb, head, voice, jaw tremor		
	Motor and phonic tics		
	Palatal myoclonus		
	Nystagmus and oscillopsia		
	Myokymia		
	Spasticity (stroke, cerebral palsy, head injury, multiple sclerosis)		
	Stuttering		
	Ophthalmologic disorders		
	Strabismus		
	Protective ptosis		
	Pain disorders		
	Headaches – chronic migraine especially		
	Tennis elbow and other sports injuries		
	Chronic anterior knee pain		
	Lumbosacral strain and back spasms		
	Radiculopathy with secondary muscle spasm		
	Myofascial pain syndromes		
	Gastrointestinal disorders		
	Sialorrhea		
	Spasm of the inferior constrictor of the pharynx (cricopharyngeal muscle)		
	Achalasia (lower esophageal sphincter spasm)		
	Obesity (distal stomach)		
	Spasm of the sphincter of Oddi		
	Anal fissure		
	Constipation		
	Anismus		
	Genito-urologic disorders		
	Spastic bladder		
	Detrusor-sphincter dyssynergia		
	Overactive bladder		
	Prostatic hypertrophy		
	Vaginismus		
	(continued)		

Table 1   (continued)	Secretory disorders
	Hyperlacrimation
	Drooling (sialorrhea)
	Hyperhidrosis
	Gustatory sweating
	Cosmetic
	Glabellar wrinkles
	Brow furrows
	Frown lines
	"Crow's feet"
	Platysma lines
	Facial asymmetry
	Acne
	Others
	Diabetic foot ulcers
	Pruritus

## 2 Mechanism of Action of BoNT

## 2.1 Different and Unique Properties of Various Serotypes of BoNT

The therapeutic value of BoNT is due to its ability to produce local paralysis when injected into a muscle by preventing the release of acetylcholine (Ach) from the presynaptic nerve terminal, also referred to as chemodenervation. There are seven immunologically distinct BoNTs (A-G), with type A being the most studied and most widely used. Synthesized as single-chain polypeptide (molecular weight of 150 kD), each toxin molecule has relatively little potency until it is cleaved by trypsin or bacterial enzymes into a heavy chain (100 kD) and a light chain (50 kD). BoNT exerts its action in a three-step process that involves binding to the receptor to the presynaptic membrane through its heavy chain, internalization of the whole toxin by endocytosis, and enzymatic action of the light chain after it has been cleaved from the heavy chain. The light chain acts as a zinc-dependent protease that selectively cleaves proteins, known as the SNARE (soluble N-ethylmaleimidesensitive factor attachment protein receptor) proteins, which are critical for fusion of intraneuronal presynaptic vesicle with the presynaptic membrane, thus preventing the release of Ach. The SNARE target proteins are different for different immunotypes of BoNT. The light chains of both BoNT-A and BoNT-E cleave SNAP-25, but at different sites. The light chains of BoNT-B, -D, and -F cleave synaptobrevin-2, also known as VAMP (vesicle-associated membrane protein), an integral protein of the synaptic vesicle membrane. BoNT-C cleaves both SNAP-25 and syntaxin, another plasma membrane-associated protein. Only types A and B are used clinically.

Generic name	Brand name(manufacturer)	Target
Onabotulinumtoxin A	Botox <sup>®</sup> (Allergan, Inc., Irvine, CA, USA) or Prosigne <sup>®</sup> (China)	SNAP 25
Abobotulinumtoxin A	Dysport <sup>®</sup> (Beaufour Ipsen, France-UK)	SNAP 25
Incobotulinumtoxin A	Xeomin <sup>®</sup> (Merz Pharmaceuticals GmbH, Frankfurt, Germany)	SNAP 25
Rimabotulinumtoxin B	Myobloc <sup>®</sup> (US) (WorldMeds, Louisville, KY, USA) or Neurobloc <sup>®</sup> (Europe)	VAMP (synaptobrevin)

Table 2 Available types of BoNT in the USA

Legend: BoNT botulinum toxin, SNAP soluble N-ethylmaleimide-sensitive factor attachment protein receptor, VAMP vesicle-associated membrane protein

There is considerable evidence that BoNT injected peripherally also influences central nervous system function. By blocking gamma as well as alpha motor neurons, there is denervation of intrafusal muscle fibers. This reduces muscle spindle afferent input to the central nervous system and thereby modifies sensorimotor and proprioceptive pathways (Giladi 1997; Hallett 2000; Rosales and Dressler 2010). These mechanisms may contribute to the therapeutic effects of BoNT in focal dystonias beyond the effects anticipated on the basis of muscle relaxation alone. There is also evidence that BoNT spreads not only to adjacent muscles but also to muscles on the contralateral side of the body (Frick et al. 2012). This is based on a study in which botulinum toxin (2.5 U) was injected into the tibialis muscle of anesthetized rats (n = 26) whereas control animals (n = 25)received a saline injection. Neuromuscular function, pharmacology, and expression of acetylcholine receptors (nAChRs) were evaluated in the tibialis at 0, 4, and 16 days after injection on the opposite side and in saline-injected controls. Although the mechanism of spread to the contralateral sides is not clear, the investigators demonstrated a decrease in specific twitch tension on the contralateral side and an increased sensitivity to atracurium on the toxin-injected side, despite upregulation of expression of acetylcholine receptors.

The original commercial preparation of BoNT was onabotulinumtoxin A, marketed as Botox<sup>®</sup> (Allergan, Inc., Irvine, CA, USA). Other forms of BoNT-A clinically available in the USA are abobotulinumtoxin A or Dysport<sup>®</sup> (Beaufour Ipsen, France-UK) and incobotulinumtoxin A or Xeomin<sup>®</sup> (Merz Pharmaceuticals GmbH, Frankfurt, Germany) (Albanese 2011). Another type of BoNT-A currently in clinical trials is PureTox<sup>®</sup> (Johnson & Johnson, USA). In addition, there is a Chinese form of BoNT-A (Prosigne or CBTX-A, Lanzhou Biological Products Institute, China), a Korean preparation (Meditoxin<sup>®</sup>/Neuronox<sup>®</sup>, Medy-Tox, South Korea), and many other formulations in experimental clinical trials and in development (Table 2).

Currently, the only preparation of BoNT-B is rimabotulinumtoxin B, known by the brand name Myobloc<sup>®</sup> (USA) or Neurobloc<sup>®</sup> (Europe) (US WorldMeds, Louisville, KY, USA). BoNT-B is an antigenically distinct form of BoNT and has unique physical and clinical properties that distinguish it from BoNT-A (Sadick 2003).

It is important to note that the biologic activity, measured in units, is different for the different products. Each preparation should be considered unique in its potency and properties and is measured with proprietary units that are considered to be non-interchangeable (Albanese 2009, 2011). However, through clinical practice when switching from one to another product, also supported by numerous comparison studies, many clinicians have used the following conversion ratios: Dysport<sup>®</sup> versus Botox<sup>®</sup> as 2.5 to 1 (Marchetti et al. 2005) and Myobloc/Neurobloc<sup>®</sup> versus Botox<sup>®</sup> as 50:1 (Pathak et al. 2006), whereas Xeomin<sup>®</sup> and Botox<sup>®</sup> seem to be equivalent in their potency (Jost et al. 2007; Frevert 2010; Dressler et al. 2011).

In Botox<sup>®</sup> and Dysport<sup>®</sup>, the BoNT component is formed by botulinum neurotoxin and by nontoxic proteins known as complexing proteins (Dressler and Benecke 2007), whereas the purification and manufacturing process for Xeomin<sup>®</sup> removes the complexing proteins (Frevert 2009b) giving it the highest specific neurotoxin activity (Frevert 2010).

## 2.2 Antigenicity and Immunoresistance

In addition to the biologically active toxin, many products include various amounts of nontoxin proteins like hemagglutinins and other proteins that are not required to stabilize the toxin but can have some immune-stimulating activity (Frevert 2009a). Xeomin<sup>®</sup> and PureTox<sup>®</sup> are free of such complexing proteins (Frevert 2009b). Some patients stop responding to BoNT injections after sometimes years of benefit. This seems to be secondary to the production of blocking antibodies directed against the heavy chain of the BoNT molecule (Jankovic and Schwartz 1995). Methods used to detect blocking antibodies include the mouse protection assay, the mouse phrenic nerve hemidiaphragm test, and many other tests (Hanna et al. 1999). The most commonly available test is the Western blot assay, but because of its lack of sensitivity and specificity, this test does not reliably predict true immunoresistance (Hanna and Jankovic 1998). A unilateral brow injection (UBI) is a useful clinical test, as inability to frown on the injected side due to weakness of the procerus and corrugator muscles confirms the absence of clinically meaningful immunoresistance (Hanna et al. 1999). The original preparation of Botox<sup>®</sup> contained 25 ng of neurotoxin complex protein per 100 U, but in 1997, the Food and Drug Administration (FDA) approved a new preparation that contains only 5 ng per 100 U, which has been associated with lower antigenicity (Jankovic et al. 2003). Depending on the technique used to detect blocking antibodies, the risk of antibodies to Botox® has markedly decreased (Mejia et al. 2005) and is now estimated to be about as low as 1.2 % of patients receiving the product repeatedly for up to 4 years (Brin et al. 2008). BoNT-B may be a useful alternative for patients who develop resistance to BoNT-A (Berman et al. 2005), but this preparation may be associated with higher risk of antigenicity (Jankovic et al. 2006). In one study, the effect of BoNT-B was sustained for up to 2.5 years in type A-resistant patients, but the magnitude of response diminished over time (Factor et al. 2005). Although the various neurotoxins are antigenically different, they contain a common subunit structure, and cross-reactive epitopes may cause cross-neutralization of antibodies (Atassi et al. 2008). Most patients who develop blocking antibodies to one type of BoNT thus have an increased risk of developing blocking antibodies to a different type of BoNT despite a possible initial good response to the new BoNT. While low antigenicity has been predicted with formulations of BoNT without complexing proteins (e.g., Xeomin<sup>®</sup> and PureTox<sup>®</sup>), no long-term data are yet available to support this notion.

Although frequently suggested, an association between a large cumulative dose of BoNT and long treatment duration with the development of anti-BoNT antibodies could not be confirmed (Bakheit et al. 2011).

## 2.3 Adverse Effects

Most of the adverse effects of BoNT are not preventable, but some are clearly associated with wrong dosage, inappropriate selection of muscles, and faulty injection techniques. Clinicians inexperienced with BoNT treatment should consider referral to a specialist experienced with proper use of BoNT. There are many controversies related to BoNT techniques, including the role of electromyography (EMG) in guiding injections, particularly to reach certain muscles (Comella et al. 1992). Although EMG guidance may be appropriate for BoNT treatment of limb dystonia, this does not mean that placement with EMG guidance correlates with better results, since the selection of the muscle involved in the hand dystonia is based on clinical examination and not on EMG (Jankovic 2001). Two class II methodological studies compared EMG versus muscle stimulation for needle localization; one of these studies showed enhanced accuracy of needle placement under EMG guidance (Molloy et al. 2002) while the other was inconclusive (Geenen et al. 1996). A more recent simple blind randomized control study showed that EMG guidance was associated with slightly better clinical outcome and a lower dose of BoNT in the treatment of cervical dystonia (CD), but the control group had a duration of symptoms twice as long as the EMG-guided group, so the results are difficult to interpret (Werdelin et al. 2011).

Side effects of BoNT injection usually result from an excessive weakness in the injected muscles. Local reactions include pain, edema, erythema, ecchymosis, headache, and short-term hyperesthesia (Lu and Lippitz 2009). The toxin can also diffuse to adjacent muscles and result in unwanted weakness. A number of factors seem to influence diffusion of BoNT including injection technique, concentration, and volume (Ramirez-Castaneda et al. 2013). Rarely, the toxin spreads systemically to distant sites resulting in mild generalized muscle weakness, malaise, nausea, fatigue, rash, or flu-like symptoms (Naumann et al. 2006; Baizabal-Carvallo et al. 2011; Dressler 2010a). It has been proposed that systemic distribution may be mediated by capillary or venous uptake and the volume of fluid used for BoNT reconstitution may increase systematic spread (Crowner et al. 2010).

The most common adverse effects encountered when using BoNT for CD include injection-site pain, muscle weakness, dysphagia, dry mouth, and flu-like symptoms (Poewe et al. 1998; Mejia et al. 2005; Naumann et al. 2006). Flu-like

symptoms occur in 1.7–20 % of BoNT-A injections (Baizabal-Carvallo et al. 2011) without marked differences between various serotypes of formulations although head-to-head comparisons are lacking. This side effect, however, seems to be more frequent with BoNT-B, observed in 5–55 % of injections (Baumann et al. 2003; Baumann et al. 2005a, b). The BoNT-related flu-like symptoms are typically mild, develop within 10 days of the injection, and last an average of 2–3 days, but may last up to 14 days and in rare cases render the patient bedridden. The symptom can be usually relieved with over-the-counter antipyretics and analgesics (Baizabal-Carvallo et al. 2011).

Results of a meta-analysis of 36 randomized controlled studies comprised of 1,425 subjects who received treatment with Botox<sup>®</sup> for various indications reported a rate of mild-to-moderate adverse effects of approximately 25 % in the Botox<sup>®</sup>-treated group compared with 15 % in the control group (p < 0.001). No severe adverse events, however, have been reported (Naumann and Jankovic 2004). In February 2008 the FDA expressed concern about potential severe adverse effects (respiratory compromise and death) associated with the use of onabotulinumtoxin A, based on submitted reports of children treated for cerebral palsy-associated limb spasticity. According to evidence-based clinical guidelines published by the American Academy of Neurology (AAN) in 2010, BoNT is considered to be an effective treatment for spasticity in children and adolescent with cerebral palsy (Delgado et al. 2010). In the 17 studies they reviewed, the most common adverse effects were localized pain, unsteadiness and increased falls, and fatigue. Although excessive weakness was noted in some cases, no deaths were reported.

The Cervical Dystonia Patient Registry for the Observation of Botulinum Toxin Type A Efficacy Study (CD-PROBE) is an ongoing registry aimed at understanding the use of Botox<sup>®</sup> in the treatment of CD (Jankovic et al. 2011a). Preliminary results from CD-PROBE suggest that repeated injections of Botox<sup>®</sup> have a low rate of adverse effects and are associated with meaningful and sustained relief of CD pain and improvement in quality of life (Charles et al. 2010). This has been confirmed by other retrospective series (Camargo et al. 2011), and there are other longitudinal observational studies currently under way designed to provide data on long-term clinical use of BoNT (Fernandez et al. 2013a, b).

Patients with post-polio syndrome and Eaton–Lambert syndrome, however, have been reported to have generalized weakness after local BoNT (Erbguth et al. 1993). Other contraindications to the use of BoNT include myasthenia gravis, motor neuron disease, concurrent use of aminoglycoside antibiotics, and pregnancy, although women, inadvertently injected during pregnancy, reported no untoward side effects to the fetus (Jankovic 2010a).

## 3 Clinical Application

BoNT is currently used in a wide variety of disorders, a few of which have been approved by the FDA (Table 3) or recommended by medical organizations (Table 4).

Trade		
name	Drug name	FDA-approved indications
Botox®	Onabotulinumtoxin A	Cervical dystonia, severe primary axillary hyperhidrosis, strabismus in patients $\geq$ 12 years old, blepharospasm in patients $\geq$ 12 years of age. Upper limb spasticity in adult patients. Temporary improvement in the appearance of moderate to severe glabellar lines
Dysport®	Abobotulinumtoxin A	Cervical dystonia, temporary improvement in the appearance of moderate to severe glabellar lines
Xeomin®	Incobotulinumtoxin A	Cervical dystonia, blepharospasm
Myobloc®	Rimabotulinumtoxin B	Cervical dystonia

Table 3 FDA-approved botulinum toxin products

Table 4	Level	of evidence	for clinical	use of BoNT
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Level A (established effective)	Blepharospasm
Level A (established effective)	Cervical dystonia
	Spasticity in adults
	1 2
	Equinus varus deformity in children with cerebral palsy
	Axillar hyperhidrosis
	Anal fissure
	Idiopathic detrusor overactivity
Level B	Hemifacial spasm
(possibly effective)	Adductor laryngeal dystonia
	Adductor spasticity in children with cerebral palsy
	Focal upper extremity dystonia
	Upper extremity essential tremor
	Palmar hyperhidrosis
	Pain control in children with cerebral palsy and upper extremity
	spasticity
	Allergic rhinitis
	Sialorrhea.
Level C	Gustatory sweating
(probably effective)	Oromandibular dystonia
·····	Laryngeal dystonia.

#### 3.1 Neurological Disorders

Since the first placebo-controlled trials of BoNT in the treatment of CD (Tsui et al. 1985, 1986) and cranial dystonia, including blepharospasm (Jankovic and Orman 1987), hundreds of other studies have been published confirming the efficacy and safety of BoNT in a variety of movement disorders and other neurologic and non-neurologic disorders.

The Therapeutics and Technology Assessment (TTA) Subcommittee of the American Academy of Neurology, in their 2008 evidence-based review (currently being revised), assigned level A recommendation for the use of BoNT for the treatment of CD and a level B recommendation for blepharospasm (Simpson et al. 2008a, b). This evidence-based review was recently updated by a panel of experts (Hallett et al. 2013). The panel evaluated published data on the four individual commercially available formulations: abobotulinumtoxinA (A/Abo), onabotulinumtoxinA (A/Ona), incobotulinumtoxinA (A/Inco), and rimabotulinumtoxinB (B/Rima) and made the following recommendations: "For the treatment of blepharospasm, the evidence supported a Level A recommendation for BoNT-A. A/Inco, and A/Ona; a Level B recommendation for A/Abo; and a Level U recommendation for B/Rima. For hemifacial spasm, the evidence supported a Level B recommendation for BoNT-A and A/Ona, a Level C recommendation for A/Abo, and a Level U recommendation for A/Inco and B/Rima. For the treatment of oromandibular dystonia, the evidence supported a Level C recommendation for BoNT-A, A/Abo, and A/Ona, and a Level U recommendation for A/Inco and B/Rima. For the treatment of cervical dystonia, the published evidence supported a Level A recommendation for all four BoNT formulations. For limb dystonia, the available evidence supported a Level B recommendation for both A/Abo and A/Ona, but no published studies were identified for A/Inco or B/Rima, resulting in a Level U recommendation for these two formulations. For adductor larvngeal dystonia, evidence supported a Level C recommendation for the use of A/Ona, but a Level U recommendation was warranted for B/Rima, A/Abo, and A/Inco. For the treatment of focal tics, a Level U recommendation was warranted at this time for all four formulations. For the treatment of tremor, the published evidence supported a level B recommendation for A/Ona, but no published studies were identified for A/Abo, A/Inco, or B/Rima, warranting a Level U recommendation for these three formulations".

#### 3.1.1 Blepharospasm

In 1987, Jankovic and Orman reported the results of a double-blind, placebocontrolled trial of onabotulinumtoxin A (Botox<sup>®</sup>) in 28 patients with cranial-cervical dystonia, including blepharospasm, oromandibular dystonia, and CD, the results of which were in part used by the FDA to approve Botox<sup>®</sup> in 1989 as a therapeutic agent in patients with strabismus, blepharospasm, and hemifacial spasm (Jankovic and Orman 1987). Subsequently there have been many other studies confirming the safety and efficacy of BoNT-A in the treatment of blepharospasm (Elston 1987; Mejia et al. 2005; Jankovic 2004; Kenney and Jankovic 2008; Gil Polo et al. 2013). Because of the paucity of double-blind, placebo-controlled trials (Roggenkämper et al. 2006), the TTA Subcommittee of the American Academy of Neurology concluded that there is only level B (probably effective) evidence for the efficacy of BoNT-A for the treatment of blepharospasm (Simpson et al. 2008b). This recommendation will likely be changed to level A as a result of more recent, well-designed, controlled trials with various BoNT products.

The onset of the Botox<sup>®</sup> effect occurs within 24–72 h, peaks at 2–4 weeks, and lasts 3–6 months (Tan 2005). While about a third of all treatment sessions are followed by some side effects (ptosis, blurring of vision or diplopia, tearing, and local hematoma), only 1–2 % affect patient's functioning, and complications

usually improve spontaneously in less than 3 weeks (Elston 1987; Roggenkämper et al. 2006). The relationship between blepharospasm, BoNT, and occurrence of dry eyes in some patients is not well understood.

In a small prospective, randomized, double-blind study on 21 patients with blepharospasm, Prosigne<sup>®</sup> was found to have similar efficacy, safety, and tolerability than Botox<sup>®</sup> with a 1:1 dose equivalence (Quagliato et al. 2010a).

Dysport<sup>®</sup> has also showed sustained efficacy and favorable safety profile with 80 U/eye (Truong et al. 2008a; Truong 2012). The onset of action occurs within 2 weeks of treatment, peaks at 4–8 weeks, and lasts 10–16 weeks (Odergren et al. 1998). In an early double-blind study in patients with blepharospasm who received one injection of Botox<sup>®</sup> and one injection of Dysport<sup>®</sup> in two separate treatment sessions, there was similar efficacy and no significant difference in the duration of the treatment effect between the two preparations, although there were significantly (p < 0.05) fewer side effects with Botox<sup>®</sup> (particularly ptosis; p < 0.01) (Nüssgens and Roggenkämper 1997). A more recent study however noted that the mean duration of improvement of blepharospasm was significantly longer for Dysport<sup>®</sup> (Bentivoglio et al. 2009), but confirmed Dysport<sup>®</sup> had a higher rate of adverse effects. Finally, another study that evaluated the effect of Botox<sup>®</sup> after switching from Dysport<sup>®</sup> (Bihari 2005).

The efficacy and safety of Xeomin<sup>®</sup> in the treatment of blepharospasm was demonstrated by a prospective, double-blind, placebo-controlled, randomized, multicenter study involving 109 patients (Jankovic et al. 2011b). An open-label extension of that study demonstrated maintained efficacy and good tolerability of Xeomin<sup>®</sup> at 48 weeks, after an average of five injections (Grafe and Hanschmann 2010; Grafe et al. 2010). In that study, the most commonly reported adverse effects were eyelid ptosis, dry eye, and dry mouth. Xeomin<sup>®</sup> was found to be clinically non-inferior to Botox<sup>®</sup> in head-to-head studies (Roggenkämper et al. 2006) (Jankovic 2009a; Wabbels et al. 2011). There is no difference between the two preparations in efficacy, safety profile, onset of action, duration, and waning of effect (Jost et al. 2007; Grafe et al. 2009a). Xeomin<sup>®</sup> is FDA approved for the treatment of blepharospasm (Barnes et al. 2010).

Myobloc<sup>®</sup> has also been used successfully in the treatment of blepharospasm, especially in Botox<sup>®</sup>-resistant patients (Dutton et al. 2006), although double-blind controlled studies in these disorders are lacking (Colosimo et al. 2003). However, side effects are more common than typically expected of BoNT-A, particularly pain on injection, ptosis, dry mouth, and dry eyes (Dutton et al. 2006).

Apraxia of eyelid opening, especially when associated with blepharospasm, seems to improve with injection of the pretarsal orbicularis oculi and injection of the pretarsal portion of the eyelid, or pars ciliaris at the lid margin seems critical for the treatment to be effective in that condition (Inoue and Rogers 2007).

#### 3.1.2 Other Cranial Dystonias

Oromandibular dystonia (OMD) is among the most challenging forms of focal dystonia to treat as it rarely improves with medications (Jankovic and Orman 1988;

Klawans and Tanner 1988; Sankhla et al. 1998). Also, there are no surgical treatments, and BoNT therapy can be complicated by swallowing problems. Clenching, trismus, and bruxism are frequent manifestations of oromandibular dystonia. Although most of the data on BoNT in the management of oromandibular dystonia (OMD) come from open-label studies, these provide compelling evidence on the efficacy of both BoNT-A and BoNT-B in patients with OMD (Wan et al. 2005; Jankovic and Orman 1987; Tan and Jankovic 2000; Laskawi and Rohrbach 2001; Alonso-Navarro et al. 2011). Patients with dystonic jaw closure, treated with injections into the masseter and temporalis muscles, generally respond better than those with jaw-opening dystonia. Although most patients with jaw-opening dystonia benefit from injections into the submentalis muscle complex, some may also require injections into the lateral ptervgoid muscles (Evidente and Adler 2010). As with blepharospasm, the improvement is usually noted within the first 5 days and persists for about 3–4 months. Early BoNT treatment of oromandibular dystonia. particularly when associated with trismus and bruxism, may prevent dental and other complications, including the temporomandibular joint syndrome.

Frowning without blepharospasm, as a manifestation of upper facial dystonia, often seen in parkinsonian patients, particularly those with progressive supranuclear palsy, can be also effectively treated with BoNT (Hirota et al. 2008).

## 3.1.3 Laryngeal Dystonia (Spasmodic Dysphonia)

Several studies have established the efficacy and safety of Botox<sup>®</sup>, Dysport<sup>®</sup>, and Myobloc<sup>®</sup> in the treatment of adductor laryngeal dystonia, produced by involuntary contraction of the thyroarytenoid muscle, and this approach is now considered by most to be the treatment of choice for spasmodic dysphonia (Adler et al. 2004b; Wan et al. 2005; Upile et al. 2009). Adverse effects include transient breathy hypophonia, hoarseness, and rarely dysphagia with aspiration (Wan et al. 2005). This approach usually requires a multidisciplinary team, consisting of an otolaryngologist experienced in laryngeal injections and a neurologist knowledgeable about motor disorders of speech and voice. There are three approaches currently used in the BoNT treatment of spasmodic dysphonia: (1) unilateral EMG-guided injection; (2) bilateral approach, injecting with EMG guidance in each vocal fold; and (3) an injection via indirect laryngoscopy without EMG. Irrespective of the technique, most investigators report about 75-95 % improvement in voice symptoms (Jankovic 2010a). However, when Dysport<sup>®</sup> is used, it seems that unilateral injections are safer than, and as effective as, bilateral injections (Troung et al. 1991; Upile et al. 2009). BoNT produces less consistent benefits when spasmodic dysphonia is accompanied by voice tremor. As for abductor spasmodic dysphonia, one class III prospective, randomized, crossover treatment study involving 15 patients showed no objective benefit of Botox<sup>®</sup>, whether injected percutaneously or transnasally into the posterior cricoarytenoid muscle (Bielamowicz et al. 2001). However, anecdotal experience from centers that frequently treat patients with spasmodic dysphonia indicates that BoNT-A is an effective treatment even in the abductor form, although the effects are more consistent in patients with adductor form of spasmodic dysphonia (Evidente and Adler 2010).

#### 3.1.4 Cervical Dystonia (CD)

The introduction of BoNT in the treatment of CD has changed the natural history of this disease in that cervical contractures are now much less frequent than they were prior to BoNT (Ramirez-Castaneda and Jankovic 2013). The efficacy and safety of BoNT in the treatment of CD have been demonstrated in several controlled and open trials (Brashear et al. 1999; Brin et al. 1999; Truong et al. 2005, 2010), and the TTA Subcommittee of the American Academy of Neurology concluded that there is level A (effective) evidence for the efficacy of Botox<sup>®</sup> for the treatment of CD (Simpson et al. 2008b). The most common treatment-related adverse events of BoNT in CD are mild-to-moderate weakness, dysphagia, dry mouth, neck pain, and injection-site pain.

In one of the first clinical trials to evaluate the efficacy of BoNT for the treatment of CD, treatment with Botox<sup>®</sup> produced significant improvement including reduction in pain. Side effects were few and no significant systemic adverse reactions were noted (Tsui et al. 1986). These results have since been confirmed in numerous short-term (Brin et al. 1987; Gelb et al. 1989; Greene et al. 1990) and long-term studies (Jankovic and Schwartz 1990; Ramirez-Castaneda and Jankovic 2013).

In a small prospective, randomized, double-blind study on 24 patients with CD, Prosigne<sup>®</sup> was found to have similar efficacy, safety, and tolerability than Botox<sup>®</sup> with a 1:1 dose equivalence (Quagliato et al. 2010b).

Dysport<sup>®</sup> was also shown to be significantly more efficacious than placebo at weeks 4, 8, and 12 in a multicenter, double-blind, randomized, placebo-controlled trial on 80 patients with blurred vision and weakness as the only side effects to occur significantly more often with Dysport<sup>®</sup> (Truong et al. 2005). These results were confirmed by multiple other randomized, double-blind, placebo-controlled trials (Truong et al. 2008b, 2010; Wissel et al. 2001; Truong and Jost 2006). Dysport<sup>®</sup> has also been shown to be more effective and better tolerated than trihexyphenidyl in the treatment of CD (Brans et al. 1996).

A recent study that evaluated the safety, efficacy, and duration of effect of Botox<sup>®</sup> after switching from Dysport<sup>®</sup> in patients with CD showed that Botox<sup>®</sup> had more efficacy and a longer duration of effect than Dysport<sup>®</sup> (Bihari 2005). However, another study showed Dysport<sup>®</sup> to be more effective than Botox<sup>®</sup> with higher adverse effects, although none of these required withdrawal of therapy or specific management (Ranoux et al. 2002). Finally, a systematic review showed Dysport<sup>®</sup> to cause more dysphagia than Botox<sup>®</sup>, possibly as a result of a greater diffusion (Chapman et al. 2007).

Myobloc<sup>®</sup> has been shown to be more effective than placebo in both type A-responsive and type A-resistant CD (Brashear et al. 1999; Brin et al. 1999; Lew et al. 1997) and may be an alternative in patients who have developed resistance to BoNT-A (Brin et al. 1999). Botox<sup>®</sup> and Myobloc<sup>®</sup> were directly compared at 1:40 dose ratio in 139 patients with CD, with no difference in improvement at 4 weeks following injection, but with more frequent dysphagia and dry mouth with Myobloc<sup>®</sup> (Comella et al. 2005). Another study on 111 patients with CD randomized to receive either Botox<sup>®</sup> or Myobloc<sup>®</sup> at a dosing ratio of 66.7 showed no significant differences in the efficacy or occurrence of

injection-site pain and dysphagia. Dry mouth however was consistently found to be more frequent with Myobloc/Neurobloc<sup>®</sup> than with Botox<sup>®</sup> (Pappert et al. 2008). A report of a 44 % of BoNT-B antibody-induced therapy failure in toxin-naïve CD patients after a relatively short exposure may however limit its use (Dressler and Bigalke 2005).

The risk of immunoresistance seems to be higher with Myobloc<sup>®</sup> than with Botox<sup>®</sup>. In a prospective, open-label, multicenter study of 333 BoNT-naïve patients with CD, after a median of 9 of Botox<sup>®</sup> treatments over a mean of 2.5 years (range: 3.2 months to 4.2 years), only 1.2 % tested positive for blocking antibodies (Brin et al. 2008). In one study of 100 patients with CD followed for 42 months over a mean of 5 (up to 12) visits, a third of the patients who were negative for BoNT-B antibodies at baseline became positive for such antibodies at the last visit (Jankovic et al. 2006). Thus, although BoNT-B offers a useful alternative to patients with immunoresistance to BoNT-A, long-term efficacy is limited by the development of blocking antibodies, probably as a result of the cross-reactivity between the two serotypes.

Xeomin<sup>®</sup> was shown to be an effective and safe treatment of CD in a doubleblind, placebo-controlled study of 233 patients, 39 % of which were BoNT naïve (Comella et al. 2011). It was found to be as effective and safe as Botox<sup>®</sup> in three different clinical trials totaling 816 patients with a 1:1 dose ratio (Jost et al. 2007) and in several other series (Benecke et al. 2005; Dressler 2009). Xeomin<sup>®</sup> seems to have similar efficacy and safety among treatment-naïve and previously treated patients (Grafe et al. 2009a, b, c; Fernandez et al. 2013a). Repeated treatments with Xeomin<sup>®</sup> seem effective and well tolerated (Grafe and Hanschmann 2010; Evidente et al. 2013). The clinical effect of Xeomin<sup>®</sup> begins within 1 week, peaks at approximately 4–6 weeks, and is sustained for about 110 days (Jost et al. 2007). The most frequently reported adverse effects of Xeomin<sup>®</sup> are mild dysphagia, neck pain, and muscle weakness (Comella et al. 2011). Xeomin<sup>®</sup> is FDA approved for the treatment of CD (Barnes et al. 2010).

## 3.1.5 Writer's Cramp, Other Limb Dystonias, and Axial Dystonia

The treatment of hand dystonia with BoNT is more challenging than the other dystonias, because there are more muscles involved in finely coordinated motor function required in the act of writing, dressing, and in various demanding tasks such as playing musical instruments and sport activities. Several open and doubleblind controlled trials have concluded that BoNT injections into selected hand and forearm muscles probably provide the most effective relief in patients with various task-specific and occupational dystonias. BoNT stands as the first-line treatment of choice for the majority of focal dystonias (Albanese et al. 2006; Benecke and Dressler 2007; Albanese et al. 2011) and seems to retain its efficacy a decade after initiation of treatment (Lungu et al. 2011). BoNT has also been used to effectively correct abnormal limb postures seen in other movement disorders such as foot dystonia and striatal hand in Parkinson's disease patients (Giladi et al. 1994; Pacchetti et al. 1995), progressive supranuclear palsy, and corticobasal degeneration (Cordivari et al. 2001; Vanek and Jankovic 2001). A marked improvement in the severity and disability was achieved with EMGguided injection of BoNT-A in 93 patients with writer's cramp, whereas primary writing tremor was little improved (Marion et al. 2003). In one study, 69 % of 84 musicians reported improvement with EMG-guided Dysport<sup>®</sup> injections, but only 36 % reported long-term benefit (Schuele et al. 2005). Dysport<sup>®</sup> was also shown to be effective in a class I randomized, double-blind, placebo-controlled trial in 40 patients with writers' cramp (Kruisdijk et al. 2007) as well as in a class II study (Contarino et al. 2007). On the other hand, Botox<sup>®</sup> improved focal hand dystonia in 80 % of patients in a class II double-blind, placebo-controlled, crossover study (Cole et al. 1995), with similar results found in another class II study (Tsui et al. 1993).

BoNT has also been used in the treatment of axial postural abnormalities secondary to axial dystonia, with variable success (Jankovic 2010b). One study reported improvement of six of nine patients with lateral axial dystonia (scoliosis) after EMG-guided injection of 500 U of Dysport<sup>®</sup> into paraspinal muscles at the level of L2–L5 on the side of the trunk flexion (Bonanni et al. 2007). Another study reported improvement of camptocormia in 9 of 11 patients who received 300–600 U of Botox<sup>®</sup> into the rectus abdominis (Azher and Jankovic 2005). In contrast, ultrasound-guided injection of the iliopsoas muscle with BoNT is not effective in camptocormia (von Coelln et al. 2008).

#### 3.1.6 Hemifacial Spasm

Hemifacial spasm is defined as a neurologic disorder manifested by involuntary, recurrent twitches of the eyelids and other muscles of only one side of the face (Yaltho and Jankovic 2011). The muscular contractions result from an irritative lesion of the ipsilateral facial nerve, most commonly from compression by a vascular loop. While microvascular decompression of the facial nerve has a high success rate, this surgical treatment is associated with certain risks, such as permanent facial paralysis, deafness, stroke, and even death. Therefore, local injections of BoNT into involved facial muscles offer a useful alternative to surgical therapy. Nearly all patients improve; the complications are minimal and transient, and the approach can be individualized by injecting only those muscles – the contractions of which are most disturbing to the patient. Along with blepharospasm, the FDA approved onabotulinumtoxin A injections for hemifacial spasm in 1989.

The average latency from injection to the onset of benefit is 5.4 days, and the total duration of benefit averages 18.4 weeks. Side effects include facial weakness, lid weakness, ptosis, teary or dry eyes, diplopia, and hematoma. The average duration of improvement in hemifacial spasm, 5 months, is longer than in any of the dystonic disorders, and rare patients have achieved long-lasting remissions. Facial myoclonus associated with Rasmussen encephalitis, similar to hemifacial spasm, but pathophysiologically related to focal cortical seizure, has been also reported to improve with BoNT (Jankovic 2010a).

The frequent use of Botox<sup>®</sup> and Dysport<sup>®</sup> in the treatment of hemifacial spasm stems mainly from extensive open-label and clinical experience (Gil Polo et al. 2013)

rather than from controlled trials (Jankovic et al. 1990; Jost and Kohl 2001; Kenney and Jankovic 2008). In one class II prospective, blinded study, Botox<sup>®</sup> was showed to be safe and effective (Yoshimura et al. 1992).

One study that evaluated the effect of Botox<sup>®</sup> after switching from Dysport<sup>®</sup> showed that Botox<sup>®</sup> was more efficacious in treating hemifacial spasm dystonia or hemifacial spasm and had a longer duration of effect than Dysport<sup>®</sup> (Bihari 2005). However, a class II single-blind, randomized, parallel-design study comparing Botox<sup>®</sup> and Dysport<sup>®</sup> at a dose ratio of 1:4 in 91 patients with hemifacial spasm or blepharospasm showed similar clinical efficacy and tolerability of both products (Sampaio et al. 1997). Xeomin<sup>®</sup> has been shown to be as effective and as safe as Botox<sup>®</sup>, with a ratio of 1:1 in 17 patients with hemifacial spasm (Dressler 2009). Prosigne<sup>®</sup> was found to have similar efficacy, safety, and tolerability than Botox<sup>®</sup> with 1:1 dose equivalence on 36 patients with hemifacial spasm (Quagliato et al. 2010a).

Myobloc<sup>®</sup> has also been used successfully in the treatment of hemifacial spasm, although double-blind controlled studies in this disorder are lacking (Colosimo et al. 2003; Trosch et al. 2007).

#### 3.1.7 Tremor

Tremor accompanies dystonia in about half of all dystonic patients, and dystonic tremor improves in some patients treated for focal dystonia with BoNT. Chemodenervation with Botox<sup>®</sup> may ameliorate not only dystonic tremor but also essential tremor involving the hands, as demonstrated by at least two doubleblind, placebo-controlled studies (Jankovic et al. 1996; Brin et al. 2001). When wrist extensor injections are avoided, weakness of finger extensors, noted in the initial studies, can be prevented. BoNT can also be considered for rest tremor associated with Parkinson's disease (PD) when antiparkinsonian treatments fail to insure satisfactory relief (Jankovic and Schwartz 1991; Diamond and Jankovic 2006a, b). In an open-label pilot study, Dysport<sup>®</sup> injection in the masseters markedly improved jaw tremor in 3 PD patients (Schneider et al. 2006).

However, essential tremor and dystonic tremor seem to respond better to BoNT than rest tremors related to PD, and further studies are needed to demonstrate efficacy of BoNT for the latter and to provide insights on how to improve the treatment protocol (Jankovic 2009b).

Although Botox<sup>®</sup> is clearly a useful treatment in patients with hand tremor, it has also been found effective in the treatment of voice tremor (Adler et al. 2004a) and head tremor (Pahwa et al. 1995). In a small, randomized, crossover trial of onabotulinumtoxin A in 23 patients with disabling tremor in 33 upper limbs related to MS, there was a significant improvement after active treatment compared with that after placebo at 6 and 12 weeks after injection in the Bain score for tremor, writing, Archimedes spiral drawing, drinking from a cup, and in the 9-hole peg test (Van Der Walt et al. 2012). There was, however, no improvement in the quality of life as measured by the Quality of Life in ET Questionnaire (QUEST). Treatment was complicated by transient weakness, noted in 42.2 % of patients treated with BTX, compared to 6.1 % of patients treated with placebo (p = 0.0005).

#### 3.1.8 Tics

Motor and phonic tics associated with Tourette syndrome typically improve with antidopaminergic drugs, but when these drugs do not adequately control the tics or are associated with troublesome side effects, BoNT injections into the affected body parts not only may provide satisfactory control of the tics but also may eliminate the premonitory urge. BoNT treatment is particularly useful in the treatment of focal motor tics and phonic tics, including coprolalia. In a placebocontrolled class II study of 18 patients with simple motor tics (Marras et al. 2001), Botox<sup>®</sup> treatment was associated with a 39 % reduction in the number of tics per minute within 2 weeks after injection against a 6 % increase in the placebo group (p = 0.004). In addition, there was a 0.46 reduction in "urge scores" with BoNT, against a 0.49 increase in the placebo group (p = 0.02). This preliminary study, however, lacked the power to show significant differences in other measured variables, such as severity score, tic suppression, pain, and patient global impression. However, it measured the results at 2 weeks, when the full effects of Botox® may have not yet been appreciated, and measured the effect of one treatment only, whereas several adjustments in doses and sites of injections over several treatment visits are usually needed in clinical practice. Another class IV open-label study of 15 patients with simple motor tics showed long-term efficacy of Botox<sup>®</sup> (Rath et al. 2010), with a permanent remission of the treated tic in three patients at 10 years' follow-up. The premonitory urge was similarly reduced.

#### 3.1.9 Spasticity and Other Hypertonic Disorders

In addition to involuntary movement disorders, BoNT has been used effectively to treat spasticity associated with cerebral palsy, multiple sclerosis, or following stroke.

There is extensive data from clinical trials indicating that Botox<sup>®</sup> is well tolerated and effective in reducing focal upper limb hypertonia following stroke (Childers et al. 2004; Gordon et al. 2004; Turkel et al. 2006; Elia et al. 2009; Simpson et al. 2009). A meta-analysis of seven multicenter, randomized, doubleblind, placebo-controlled, parallel-group trials totaling 544 poststroke patients showed that Botox<sup>®</sup> decreased poststroke spasticity in upper limb-injected muscles proportionally to the dose of BoNT used. Injected muscles included the flexor carpi ulnaris (FCU), flexor carpi radialis (FCR), flexor digitorum superficialis (FDS), flexor digitorum profundus (FDP), and biceps brachii (BB) (Yablon et al. 2011). A multicenter, randomized, double-blind, placebo-controlled study of 120 poststroke patients with lower limb spasticity showed that a onetime Botox<sup>®</sup> injection yielded significant improvement at 4, 5, and 8 weeks (Kaji et al. 2010). Experienced injectors have recommended the following doses of Botox in the treatment of spasticity: for the lower limb, 3-6 U/kg per muscle; for the upper limb above the elbow, 2-3 U/kg per muscle; and for the upper limb below the elbow and posterior tibialis, 0.5–2 U/kg per muscle (Jankovic 2010a).

A double-blind, placebo-controlled, crossover study also demonstrated improvement in spasticity of thigh adductors in patients with multiple sclerosis after receiving Botox<sup>®</sup> with subsequent functional gain, specifically easier nursing care, and better comfort when sitting in a wheelchair (Snow et al. 1990). Dysport<sup>®</sup> also seems to improve upper limb and, to a lesser extent, lower limb spasticity secondary to stroke (Keam et al. 2011).

Xeomin<sup>®</sup> was also shown to be effective in reducing poststroke upper limb spasticity in a recent open-label study of 145 patients (Kanovsky et al. 2011).

Based on the review of 11 class I trials for adult upper extremity spasticity, one of them using BoNT-B, and three class I trials for lower limb spasticity, the AAN recommends that BoNT should be offered as a treatment option to reduce muscle tone and improve passive function in adults with spasticity (Simpson et al. 2008a).

Several reviews have assessed the therapeutic effect of BoNT-A, Botox<sup>®</sup> and Dysport<sup>®</sup> confounded, for spasticity of the upper or lower limbs secondary to cerebral palsy and the consequent impact on quality of life (Lannin et al. 2006; Naumann et al. 2006; Park and Rha 2006; Coutinho et al. 2011). A meta-analysis of 20 randomized controlled trials of Botox<sup>®</sup> and/or Dysport<sup>®</sup> versus placebo or rehabilitation, totaling 882 patients, showed that BoNT-A has a good safety profile during the first months of use, with however more systemic side effects. BoNT-A use was related to respiratory tract infection, bronchitis, pharyngitis, asthma, muscle weakness, urinary incontinence, falls, seizures, fever, and unspecified pain (Albavera-Hernández et al. 2009). Since a dose–response relationship has been observed in the frequency and severity of adverse events of BoNT-A (Naumann et al. 2006), it is possible that children with cerebral palsy have a higher probability of occurrence since they require high doses of toxin relative to their body mass.

In children with cerebral palsy,  $Botox^{\mathbb{R}}$  (2–8, up to 16 U per kg body weight per muscle) is often found effective when injected in calf muscles to correct equinus deformity and toe walking and in the hamstring muscles to correct crouch and scissor gait, improve sitting and hygiene care, and reduce pain. In a double-blind, controlled study, patients with cerebral palsy receiving 40-80 U of Botox<sup>®</sup> per muscle experienced a significantly greater improvement in Ashworth score and gait measures than the "low-dose" group (20-40 U per muscle) (Wissel et al. 1999). Multiple studies of Botox<sup>®</sup> or Dysport<sup>®</sup> injections into the gastrocnemius showed gait improvement over 1-3 months (Simpson et al. 2008a). One randomized, double-blind, placebo-controlled study injecting Dysport<sup>®</sup> into the thigh adductors and medial hamstrings of children with adductor spasticity showed significant improvement in knee-to-knee distance as well as adductor muscle tone (Mall et al. 2006). The results of many of the studies are difficult to interpret and compare, because patients with various forms and etiologies of spasticity were enrolled and different methodologies were used to inject and rate the patients. Although some double-blind trials have demonstrated meaningful functional improvement in patients with spasticity, other controlled studies have failed to demonstrate improvement. Many of the published studies suffer methodological problems, particularly in selecting the appropriate outcome measures that may or may not capture the functional goals of the patients. The AAN granted BoNT injections a level A of evidence for the treatment of equinovarus deformity in children with cerebral palsy and a level B for treatment of adductor spasticity and for pain control and in children with cerebral palsy and upper extremity spasticity (Level B) (Simpson et al. 2008a).

Xeomin<sup>®</sup> was reported useful in treating acquired brain injury-induced upper limb spasticity in a series of 192 patients (Barnes et al. 2010) and lower limb spasticity in one case report (Lippert-Gruner and Svestkova 2011). It has been shown to be as effective and as safe as Botox, with a ratio of 1:1 in a series of 94 patients with spasticity (Dressler 2009).

### 3.2 Painful Disorders

## 3.2.1 Headache

No significant difference between Botox<sup>®</sup> and placebo could be demonstrated in four different studies on patients suffering from episodic migraines, defined as less than 15 days of headache per month (Silberstein et al. 2000; Elkind et al. 2006; Relja et al. 2007; Saper et al. 2007; Naumann et al. 2008). Similarly, out of 4 studies of Botox<sup>®</sup> for chronic daily headache, none demonstrated a significant effect of Botox<sup>®</sup> on the primary outcome measure (Mathew et al. 2005). Botox<sup>®</sup> was also ineffective in treating chronic tension headaches in four randomized, placebo-controlled studies (Naumann et al. 2008). Another study reported that imploding headaches (head seems to be crushed, clamped, or stubbed by external forces) and ocular headaches (buildup of pressure inside the head) (Jakubowski et al. 2006).

One controlled study demonstrated the effectiveness of fixed-site administration of 100 U of Botox<sup>®</sup> in the treatment of patients with chronic migraine who specifically did not overuse pain medication (Freitag et al. 2008). However, 150 U of Botox would be needed in patients with chronic migraine and medication overuse (Grazzi, 2013). Chronic migraine is defined as headache occurring >15days per month for >3 months, with headaches occurring on >8 days being classified as migraine headaches or headaches that respond to migraine-specific medications (Olesen et al. 2006). More recently, pivotal results from the Phase III Research Evaluating Migraine Prophylaxis Therapy (PREEMPT) clinical program involving two double-blinded, placebo-controlled studies established Botox<sup>®</sup> as a safe, well-tolerated, and effective headache prophylactic treatment for chronic migraine using a combination of fixed-site administration and follow-the-pain approach, at doses ranging from 155 to 195 U administered across seven head and neck muscles every 12 weeks for up to five treatment cycles (Aurora et al. 2010; Blumenfeld et al. 2010; Diener et al. 2010; Dodick et al. 2010). However, these results were not replicated with Dysport<sup>®</sup> (Chankrachang et al. 2011).

The antinociceptive effect of BoNT in headache was initially thought to be a result of relief of muscle spasms, but in vitro studies have shown that BoNT blocks the peripheral release of pain and inflammatory neurotransmitters such as glutamate, substance P, and calcitonin gene-related peptide (Aoki 2003, 2005; Gazerani et al. 2009). Moreover, it has been inferred from animal studies that BoNT-A exerts its antinociceptive effects through retrograde transport and involvement of the central nervous system (Bach-Rojecky and Lacković 2009).

#### 3.2.2 Chronic Knee Pain

In one randomized, double-blind, placebo-controlled, crossover trial on 24 patients with chronic anterior knee pain associated with quadriceps muscle imbalance, one 500 U injection of Dysport<sup>®</sup> in the vastus lateralis associated with retraining of the vastus medialis provided marked pain improvement compared to placebo. This intervention permitted patients to restore a more balanced knee extensor control during functional activity (Singer et al. 2011).

## 3.2.3 Painful Limbs/Moving Extremities

The syndrome of painful limbs/moving extremities describes a clinical condition characterized by involuntary toe or finger movement associated with pain in the affected hand/arm or foot/leg. Botox<sup>®</sup> injections in the involved muscles have been shown to improve pain and movement in some case reports (Singer and Papapetropoulos 2007; Eisa et al. 2008).

### 3.2.4 Other Pain Syndromes

BoNT-A seems effective in controlling musculoskeletal pain, such as myofascial pain, low back pain, trigeminal neuralgia, and other chronic pain syndromes (Reilich et al. 2004; Sycha et al. 2004; Bhidayasiri and Truong 2005; Bohluli et al. 2011). However, these conditions represent a diverse group, and the results with BoNT-A have not been universally positive (Waseem et al. 2011; Zhang et al. 2011). A recent meta-analysis failed to confirm any significant improvement of neck pain after BoNT-A injection (Langevin et al. 2011a, b).

## 3.3 Hyperhidrosis

Several well-designed trials have demonstrated efficacy of intradermal injection of Botox<sup>®</sup> and Dysport<sup>®</sup> in axillar (Heckmann et al. 2001; Naumann and Lowe 2001; Doft et al. 2011; Dressler and Adib Saberi 2013) and palmar hyperhidrosis (Lowe et al. 2002; Solish et al. 2007; Ito et al. 2011). A within-subject comparison showed similar effectiveness and tolerance but longer duration of action of Dysport<sup>®</sup> compared to Botox<sup>®</sup> (Simonetta et al. 2003). Xeomin<sup>®</sup> has been shown to be as effective and as safe as Botox<sup>®</sup>, with a ratio of 1:1 in 64 patients with hyperhidrosis (Dressler 2009). This was confirmed by a double-blinded study on 46 patients complaining of axillar hyperhidrosis with injection of Botox<sup>®</sup> under one arm and Xeomin<sup>®</sup> under the other (Dressler 2010b).

Myobloc<sup>®</sup> has also been shown to improve axillary as well as palmar hyperhidrosis (Baumann et al. 2005a, b) and appears to be as effective as Botox<sup>®</sup> in the treatment of axillary hyperhidrosis at a conversion factor of 20:1, with however more discomfort on injection (Dressler et al. 2002). In a recent within-subject comparison of 10 patients, Myobloc<sup>®</sup> was significantly more effective and had a longer duration than Botox<sup>®</sup>, at a conversion ratio of 50:1 (Frasson et al. 2011). Xeomin<sup>®</sup> has also been used successfully for the treatment of axillar hyperhidrosis (Dressler 2012). The TTA Subcommittee of the AAN concluded that there is level

A evidence for the recommendation that BoNT should be offered as a treatment option for axillary hyperhidrosis and level B for palmar hyperhidrosis (Naumann et al. 2008).

Gustatory sweating is a unique entity characterized by profuse sweating of the face, scalp, and neck during or immediately after ingestion of food or drink (Naumann et al. 1997). It most often occurs as a complication of surgery to the area of the face near the parotid glands (Nolte et al. 2004). The effect of Botox<sup>®</sup> on gustatory sweating has been evaluated in one class II and three class III studies totaling 131 patients, with an effect lasting up to a year (Naumann et al. 1997; Laccourreye et al. 1999; Eckardt and Kuettner 2003; Nolte et al. 2004). At present available evidence supports a Level C recommendation for BoNT for the treatment of gustatory sweating.

## 3.4 Gastrointestinal Conditions

#### 3.4.1 Sialorrhea

Several studies, including randomized placebo-controlled trials (Ondo et al. 2004; Lagalla et al. 2006; Jackson et al. 2009), have provided evidence that Botox<sup>®</sup>, Dysport<sup>®</sup>, or Myobloc<sup>®</sup> injections in the parotid glands with or without injection in the submandibular glands may be the treatment of choice for sialorrhea and drooling associated with PD and other disorders with bulbar dysfunction, such as amyotrophic lateral sclerosis, cerebral palsy, posttraumatic encephalopathy, and bilateral strokes (Nóbrega et al. 2007; Santamato et al. 2008; Molloy 2007; Basciani et al. 2011; Chinnapongse et al. 2011; Moller et al. 2011). Ultrasound-guided injection may improve the efficacy of BoNT treatment for sialorrhea (Dogu et al. 2004). Xeomin<sup>®</sup> has been shown to be as effective and as safe as Botox<sup>®</sup>, with a ratio of 1:1 in a small series of patients with sialorrhea (Dressler 2009).

An open-label study of 30 children with neurological disorders and sialorrhea showed no difference in efficacy or side effects between BoNT-A and BoNT-B (Wilken et al. 2008). Other studies provided evidence that BoNT-B may be more effective than BoNT-A in treating sialorrhea as dry mouth is significantly more frequently noted in patients treated for CD with BoNT-B as compared to BoNT-A (Tintner et al. 2005; Jankovic 2009b; Guidubaldi et al. 2011). However, a recent randomized double-blind crossover trial demonstrated similar efficacy but shorter latency and lower cost of Myobloc<sup>®</sup> compared to Dysport<sup>®</sup> (Guidubaldi et al. 2011).

The potential adverse effects of using BoNT for sialorrhea include transient dysphagia and xerostomia although dysphagia has not been reported in any study. In fact, a study of oropharyngeal swallowing dynamics, based on swallowing videofluoroscopy, showed no difference before or 30 days after injections of BoNT-A into the parotids in patients with PD (Nóbrega et al. 2009).

## 3.4.2 Achalasia

Achalasia is manifested by spontaneous and repetitive contractions of the proximal esophagus with failure of the lower esophageal sphincter to relax during swallowing. This leads to dysphagia. Several small studies have demonstrated the beneficial effects of Botox<sup>®</sup> injections into the lower esophageal sphincter in the treatment of

achalasia, particularly in patients who are not candidates for surgery or balloon dilatation (Pohl and Tutuian 2007; Barnes et al. 2011; Lakhtakia et al. 2011).

#### 3.4.3 Constipation and Fecal Incontinence

Anismus (constipation due to functional obstruction at the pelvic outlet by paradoxical contraction of the striated sphincter muscles during defecation straining) has been suggested to represent a form of focal dystonia and has been considered as a possible etiology of constipation in some patients (Mathers et al. 1988). Several small open-label studies demonstrated the beneficial effects of Botox<sup>®</sup> anal injections in the treatment of the subtype of constipation secondary to anismus (Albanese et al. 2003; Cadeddu et al. 2005; Hompes et al. 2012). The puborectalis muscle is usually injected under transrectal ultrasonographic guidance.

On the other hand, rectal hypercontractility can lead to fecal incontinence. Intrarectal Dysport<sup>®</sup> injections have shown some improvements in fecal incontinence in a small series (Bridoux et al. 2011).

#### 3.4.4 Anal Fissure

Botox<sup>®</sup> and Dysport<sup>®</sup> injections into the internal anal sphincter seem effective in treating chronic anal fissure with a low rate of transient incontinence secondary to diffusion to the external anal sphincter and without permanent complications. However, the therapeutic response is heterogenous (Brisinda et al. 2002; Maria et al. 2000; Yiannakopoulou 2012).

## 3.5 Overactive Bladder and Other Urologic Problems

An increasing number of reports provide evidence that Botox<sup>®</sup> injections into the bladder wall is an effective treatment in increasing the bladder capacity and improving urge and incontinence in patients with overactive bladder associated with neurogenic and idiopathic detrusor overactivity (Patel et al. 2006; Chen and Liao 2011; Deffontaines-Rufin et al. 2011). However, the rate of treatment discontinuation may be as high as 61.3 % at 3 years, secondary the need to intermittent self-catheterization or urinary tract infections (Mohee et al. 2013). A randomized controlled trial comparing anticholinergic versus BoNT for urinary urge incontinence has recently started (Visco et al. 2012). The TTA Subcommittee of the AAN concluded that there is level A evidence for the recommendation that BoNT should be offered as a treatment option for this urinary disorder (Naumann et al. 2008).

Other genitourinary indications for Botox<sup>®</sup> treatment include voiding dysfunction due to benign prostatic hypertrophy (Chuang and Chancellor 2006; Marchal et al. 2011).

### 3.6 Allergic Rhinitis

Two class II placebo-controlled studies with a total of 73 patients demonstrated the efficacy of 20–30 units of Botox<sup>®</sup> in each nasal cavity to manage the symptoms of

allergic rhinitis (Unal et al. 2003; Yang et al. 2008). Botox<sup>®</sup> was also superior when compared to intranasal triamcinolone (Yang et al. 2008). However, the follow-up was only for 8 weeks, so the durability of effectiveness is unclear.

## 3.7 Cosmetic

It is beyond the scope of this review to discuss the vast literature on cosmetic use of BoNT, which represents about half of all the uses of BoNT. Botox<sup>®</sup> and Myobloc<sup>®</sup> have been used in brow lifts and for treating glabellar lines, horizontal forehead lines, crow's feet, hypertrophic orbicularis, upper nasalis (bunny) lines, repeated nasal flares, melomental folds, perioral rhytides or smoker's lines, mouth frown, peau d'orange skin, mental crease, facial asymmetry with both hyper- and hypofunctional muscle imbalance, necklace lines, and platysmal bands (Glogau 2002; Baumann et al. 2003; Choi et al. 2013a, b; Dessy et al. 2011). Overly aggressive or imprecise treatment in the cervical area can lead to incompetent mouth, weakness of the neck flexors, and dysphagia (Klein 2001). The duration of effect of BoNT for cosmetic indications ranges from 3 to 4 months and up to 6–8 months, depending on the treatment site and injection technique (Carruthers and Carruthers 2001). Myobloc® seems to have a faster onset of action by 1-2 days for all these muscle groups, but a shorter duration and lesser potency of action when compared to Botox® at a conversion factor of 1:20 to 1:100. In addition, intradermal injections with Myobloc<sup>®</sup> are more painful than with  $Botox^{\mathbb{R}}$  (Yamauchi and Lowe 2004).

Botox<sup>®</sup> injected peri-nasally at a dose of 20–25 units for facial tics also improved acne in two patients despite previous failure of several anti-acne agents. Clearing of peri-nasal acne began 1–2 weeks after each treatment and lasted 4–5 months (Diamond and Jankovic 2006a, b). Xeomin<sup>®</sup> and Dysport<sup>®</sup> have recently demonstrated efficacy in the treatment of crow's feet (Prager et al. 2011).

## 4 Conclusion

Since first recognized as the cause of food-borne botulism in the early nineteenth century, BoNT was suggested as a potential treatment for involuntary spasms and movements. Almost 200 years later, its known clinical applications span across numerous neurological and other diseases. The Neurotoxin Institute was established in 2002 as an umbrella organization providing unbiased educational programs and other information about the broad use of BoNT in different special-ties (www.neurotoxininstitute.com).

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# Neurotoxic Effects, Mechanisms, and Outcome of 192-IgG Saporin

Laura Petrosini, P. De Bartolo, and D. Cutuli

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#### Abstract

The first type-selective anti-neuronal active in vivo immunotoxin is 192 IgGsaporin. 192 IgG-saporin selectively destroys cholinergic neurons of basal forebrain that provide cholinergic input to the hippocampus, entire cortical mantle, amygdala, and olfactory bulb. Immunotoxic lesions by 192 IgG-saporin represent a valid animal model of Alzheimer's disease, given the degeneration of basal cholinergic system present in this pathology.

L. Petrosini (🖂) • P. De Bartolo

Department of Psychology, Sapienza University of Rome, Rome, Italy

I.R.C.C.S. Santa Lucia Foundation, Rome, Italy e-mail: laura.petrosini@uniroma1.it

D. Cutuli

I.R.C.C.S. Santa Lucia Foundation, Rome, Italy e-mail: laura.petrosini@uniroma1.it

Selective lesioning of cholinergic innervation by means of intracerebroventricular (i.c.v.) or intraparenchymal (i.pr.) 192 IgG-saporin is able to interfere with experience-dependent plasticity. A number of studies have demonstrated alteration of several structural and biochemical parameters related with neuroplasticity (dendritic spines and branching of pyramidal neurons, adult neurogenesis, levels of neurotrophic factors) in both cortical mantle and hippocampus.

Furthermore, lesions of the cholinergic basal forebrain affect cognitive functions, such as learning, memory, and attention, as well as sleep-waking cycle. The effects of selective immunotoxic lesions have been examined in a variety of behavioral paradigms of learning and memory. The general framework has to take into account the route of injection (i.c.v. or i.pr.), lesion extent, age of lesioning, and kind of behavior analyzed. Namely, cholinergic depletion can elicit specific learning and memory impairments as well as deficits in attentional and discriminative abilities. However, 192 IgG-saporin lesions result in overt behavioral deficits only using high demanding tasks and following high-grade CBF lesions, indicating that the relationship between CBF lesion extent and cognitive impairment is a threshold relationship in which a high degree of neuronal loss can be tolerated without detectable consequences.

#### Keywords

192 IgG-saporin • Basal forebrain cholinergic system • Cognitive functions • Neurodegenerative disorders • Neuroplasticity

List of Abb	reviations
ACh	Acetylcholine
AChE	Acetylcholine esterase
AD	Alzheimer's disease
AF64A	Aziridinium ion of ethylcholine mustard
AMPA	Aminomethylphosphonate
APP	Amyloid precursor protein
BDNF	Brain-derived neurotrophic factor
CBF	Cholinergic basal forebrain
ChAT	Choline acetyltransferase
DBBv	Diagonal band of Broca
DMTP	Delayed matching to position
DMTS	Delayed matching to sample
DNMTP	Delayed nonmatching to position
GABA	Gamma-aminobutyric acid
HAChT	High-affinity choline transport
i.c.v.	Intracerebroventricular
i.pr.	Intraparenchymal
MS	Medial septal nucleus
MWM	Morris Water Maze

# List of Abbreviat

NBM	Nucleus basalis magnocellularis
NGF	Nerve growth factor
NMDA	<i>N</i> -methyl-D-aspartate
OF	Open field
p75	$p75^{NGFFR}$ , nerve growth factor receptor
PA	Passive avoidance
RAM	Radial arm maze
RIPs	Ribosome-inactivating proteins
SI	Substantia innominata

# 1 Introduction

The cholinergic basal forebrain (CBF) is one of the diffusely projecting systems of the brain. It provides cholinergic input to the hippocampus, entire cortical mantle, amygdala, and olfactory bulb. The physiological function of the CBF is thought to modulate the excitability of cortical and hippocampal neurons. Anatomical and physiological data on the CBF encourage the notion that it modulates some cognitive functions, as learning, memory, and attention, as well as sleep-waking cycle. The evidence for this hypothesis has come from observations that muscarinic antagonists generally impair cognitive function in humans and animals, and cholinomimetics can reverse such impairments or enhance cognition when given alone (Hagan and Morris 1988). Although the precise site within the CNS for the cholinergic drug effects on behavior is uncertain, the CBF is regarded as the probable anatomical substrate for the cognitive effects of cholinergic drugs (Markowska et al. 1995). The discovery that the CBF degenerates in Alzheimer's disease (AD) and that its degree of degeneration correlates with the degree of dementia (Perry et al. 1978) has strongly encouraged interest on the role of the CBF in cognition. Degeneration of the CBF in AD has motivated investigators to attempt to model these pathological features through a variety of experimental lesioning techniques. The approaches employed have included fimbria-fornix transaction (Matsuoka et al. 1991), injection of the toxin aziridinium ion of ethylcholine mustard (AF64A) into the basal forebrain or target areas of cholinergic neurons (Walsh and Opello 1994), and intracerebral injections of excitotoxins (selective toxins for excitatory neurons, such as kainic, ibotenic, N-methyl-D-aspartate (NMDA), and quisqualic acids) (Dunnett et al. 1991). The major shortcoming of these works is that each of these techniques lacks selectivity. For example, excitotoxins are able to destroy cell soma without damaging crossing fibers. Yet, their selectivity depends on the injection site: there is great selectivity where the neurons are organized in distinct groups, as for *locus coeruleus.* Differently, since cholinergic neurons of basal forebrain are highly widespread among other cell types, the excitotoxic lesions induce unwanted loss of non-cholinergic neighboring neurons (Dunnett et al. 1991) rendering thus difficult to ascribe the eventual resulting deficits specifically to destruction of the CBF.

In the 1990s, the problem of selective lesions of the CBF has been addressed through the development of the new lesioning techniques that have proved very significant for functional neuroscience research. The power and validity of this approach reside in the nature of the lesion.

#### 2 Immunolesioning Techniques

Immunolesions use monoclonal anti-neuronal antibodies to deliver similar cytotoxin selectivity to only those neurons bearing surface moieties recognized by the antibodies. Immunolesioning can destroy a specific type of neuron locally, or after axonal suicide transport from the injection site, thus combining selectivity for both anatomical region and cell type.

Immunotoxins are composed of monoclonal antibodies conjugated to various ribosome-inactivating proteins (RIPs). The monoclonal antibody component of an immunotoxin binds to specific cell surface molecules to result in endocytosis of the antibody and its coupled RIP. Once internalized, the RIP component inhibits protein synthesis, catalytically inactivating ribosomes and resulting in cell death. If immunotoxin binds to nerve terminals or axons, it is internalized by receptor-mediated endocytosis and then travels by retrograde axonal transport to neuronal cell bodies, where the RIP acts to inhibit protein synthesis. If antigen-bearing soma is present at the injection site, then axonal transport is obviously not necessary to deliver the toxin. However, high local toxin concentrations at the injection site may produce nonspecific cytotoxicity with necrosis of all cell type.

The first immunotoxin that was reported to be effective in making neuronal lesions is OX7-saporin that consisted of the monoclonal antibody OX7 coupled to saporin. Saporin is one RIP often used; derived from the plant *Saponaria officinalis* (soapwort). OX7 recognizes Thy-1, an abundant surface molecule present on all neurons, so the resulting immunotoxin is not selective for any one type of neuron. For a review see Wiley (1992).

# 3 192 IgG-Saporin

The first type-selective anti-neuronal active in vivo immunotoxin is the 192 IgG-saporin. 192 IgG-saporin consists of the monoclonal antibody 192 IgG which is disulfide coupled to saporin. The antibody component is directed against low-affinity rat nerve growth factor (NGF) receptor (p75<sup>NGFFR</sup>). Targeting p75 serves as an ideal way to selectively lesion the CBF neurons because in the adult rat basal forebrain, only the cholinergic neurons express p75. They possess p75 NGF receptors while neurons containing other neurotransmitters in the region, and even the nearby striatal cholinergic interneurons, do not express detectable levels of same receptors. NGF plays a role in the maintenance of function of the magnocellular CBF neurons, which have NGF receptors (Thomas et al. 1991). When injected, 192 IgG-saporin binds to the surface of p75-bearing neurons and is internalized by

endocytosis. Once in the cytoplasm, the saporin moiety escapes endosomes and enzymatically inactivates the large ribosomal subunit halting protein synthesis and leading ultimately to cell death. For a review see Wiley (1992).

#### 4 Organization of the Basal Forebrain Cholinergic System

Neurons in the basal forebrain provide the major source of cholinergic innervation for the entire neocortex, hippocampus, amygdala, and olfactory bulb. CBF is a continuum of several overlapping nuclei with divergent efferent pathways, such that a single projection field is not exclusive of a particular nucleus. The forebrain and upper brainstem of the rat contain six groups of cholinergic projection neurons designated by Mesulam as Ch1-Ch6 on the basis of cytoarchitectonic criteria and connectivity patterns. The Ch terminology was originally described on the monkey brain and later proposed in extended form in the rat. The Ch1-Ch2 sectors are contained within the medial septal (MS) nucleus and the vertical limb nucleus of the diagonal band of Broca (DBBv), respectively. They provide the major cholinergic projection to the hippocampal formation. The Ch3 sector is mostly contained within the lateral part of the horizontal limb nucleus of the diagonal band and provides the primary cholinergic projection to the olfactory bulb. The Ch4 sector contains the cholinergic neurons of the nucleus basalis magnocellularis (NBM), the substantia innominata (SI), and probably also laterally situated neurons of the vertical limb nucleus as well as medially situated cells of the horizontal limb nucleus. One unifying feature of Ch4 is that its components provide the major cholinergic projection to neocortical targets, as well as to the amygdala. The Ch5-Ch6 regions are located in the pontomesencephalic reticular formation and provide the major cholinergic innervation to the thalamus. For a review see Mesulam et al. (1983).

Developmental studies using biochemical, histochemical, and immunocytochemical techniques have shown that cholinergic neurons in the rodent basal forebrain are generated prenatally according to a distinct spatiotemporal pattern and undergo extensive maturation during the first postnatal month. Cholinergic fibers from Ch1–2 or Ch4 sectors, which start growing as early as embryonic day 15, reach the appropriate target fields in the hippocampus or neocortex at about the time of birth and establish adult-like patterns of innervation in either region within the fourth postnatal week, when the postnatal maturation of cholinergic markers is fairly complete (Leanza et al. 1996).

Several studies addressing the role of cholinergic neurotransmission investigated how to use 192 IgG-saporin, evaluating extent, time course, specificity, and mechanism of action of intraparenchymal or intraventricular injections in the adult CNS.

# 5 Selectivity and Effectiveness of 192 IgG-Saporin

The selectivity and effectiveness of 192 IgG-saporin for producing lesions of the CBF has now been shown by several research groups. Both intracerebroventricular (i.c.v.)

administration of the 192 IgG-saporin and direct injection into distinctive basal forebrain nuclei induce nearly complete and specific lesions of cholinergic cells but sparing other neuronal systems in the basal forebrain.

192 IgG-saporin i.c.v. injection affects CBF as well as Purkinje cells of the cerebellum (Berger-Sweeney et al. 1994; Heckers et al. 1994) because a subset of these cells starts expressing p75 receptors of NGF during development and continues during adulthood (Pioro and Cuello 1988). Differently, 192 IgG-saporin i.c.v. injections spare the cholinergic neurons of other brain regions, the gamma-aminobutyric acidergic (GABAergic) neurons expressing parvalbumin that are intermingled within the CBF, the calbindin D-28K expressing neurons of the NBM, and the NADPH diaphorase-expressing neurons of the NBM (Heckers et al. 1994; Wiley et al. 1995). 192 IgG-saporin i.c.v. injections also result in loss of cholinergic markers such as choline acetyltransferase (ChAT) activity, acetyl-choline esterase (AChE) positive fibers, and high-affinity choline transport (HAChT) in the target fields of the CBF (Wrenn and Wiley 1998). 192 IgG-saporin i.c.v. injections at different time points of neonatal age (from 0 to 30 postnatal day) are used for studying the development of the central cholinergic system (Berger-Sweeney 1998; Leanza et al. 1996).

When injected intraparenchymally (i.pr.) into basal forebrain, 192 IgG-saporin effects are present mainly in the proximity of the injection site, sparing interspersed non-cholinergic neurons (Berger-Sweeney et al. 1994). Notably, i.pr. injections induce a more pronounced gliosis than i.c.v. injections. Delayed loss of cholinergic striatal interneurons after injection into the Ch4 sector has been described (Heckers et al. 1994); however this finding has not been replicated. Striatal cholinergic neurons express the p75 NGF receptor in early development phases, but this expression decreases drastically during adulthood (Pioro and Cuello 1988). The described loss of striatal cholinergic neurons might be due to the lesion-induced reexpression of p75 NGF receptors in the striatum (Heckers et al. 1994). The sphere of CBF-killed cells that results from i.pr. injections can be expanded by increasing the dose. However, at high dose, nonselectivity may become a problem (Wrenn and Wiley, unpublished observation). 192 IgG-saporin i.pr. injections can also be performed in the target fields (hippocampus, cortex) of the CBF. Injections into the hippocampus result in a mainly ipsilateral loss of the Ch1-Ch2 sector cholinergic neurons, an effect blocked by simultaneous injections of colchicines. Similarly, intracortical injections cause a loss of cholinergic fiber density in the cortex and a loss of Ch4 cholinergic neurons (Ohtake et al. 1997).

The selectivity and lethality of 192 IgG-saporin for CBF neurons were further demonstrated by Book et al. (1994) by using a double-labeling approach. In this study, the persistent retrograde tracer Fluoro-Gold was cortically injected 1 week before the i.c.v. injections of 192 IgG-saporin or saline. The Fluoro-Gold pre-labeled both cholinergic and non-cholinergic basal forebrain neurons. After depleting the cholinergic neurons, Ch4 was stained immunohistochemically for ChAT. These authors found that only the neurons that were double-labeled for both Fluoro-Gold and ChAT were lost in the 192 IgG-saporin-treated rats. Thus, 192 IgG-saporin selectively killed the cholinergic neurons of the Ch4 sectors, leaving intact non-cholinergic neurons.

It is important to emphasize that much of the CBF projection to amygdala is not lesioned by i.c.v or i.pr. injections of 192 IgG-saporin. Most cholinergic neurons projecting to the amygdala do not express p75 receptors; therefore, they are unaffected by 192 IgG-saporin (Heckers et al. 1994). The sparing of the cholinergic innervation to the amygdala renders the experimental immunotoxic model significantly different from the AD pathology, representing a methodological limit that has to be taken into account.

In summary, 192 IgG-saporin is the most effective and selective means for producing lesions of the rat CBF. Its advantage over previous techniques is in its ability to selectively kill cholinergic neurons of the basal forebrain and to leave intact intermingled non-cholinergic neurons. Significant drawbacks of 192 IgG-saporin include sparing of the cholinergic innervations to amygdala and destruction of some cerebellar Purkinje neurons.

## 6 Structural and Biochemical Effects of 192 IgG-Saporin

One of the most interesting functional correlates of cortical cholinergic input is its role in mediating neuroplasticity. Neuroplasticity is a lifelong process that mediates the structural and functional reaction of dendrites, axons, and synapses to experience, aging, and injury.

The selective lesioning of cortical cholinergic innervation by means of 192 IgGsaporin interferes with experience-dependent plasticity in the barrel fields. When all whiskers except for D2 and D3 were trimmed, a pairing between the D2 and D3 barrel fields occurred so the D2 neurons in the cerebral cortex started to show a greater responsivity to the stimulation of D3 than to stimulation of adjacent D1, which had been trimmed. This pairing, indicative of experience-induced synaptic plasticity, is not obtained in rats with selective immunotoxic lesions in the Ch4 (Baskerville et al. 1997). In newborn rat pups, barrel representing intact whiskers failed to show the expected expansion into the territory representing trimmed whiskers in animals with Ch4 lesions (Kilgard and Merzenich 1998). Furthermore, pairing auditory stimuli with electrical stimulation of the Ch4 in adult rats caused a long-lasting reorganization of primary auditory cortex so that the area optimally responsive to the paired tone expanded substantially. This plasticity was not observed following the selective immunotoxic destruction of cholinergic Ch4 neurons. Thus, it appears that cortical cholinergic denervation can undermine the learning-dependent reorganization of cortical representations and perhaps also the ability of the brain to keep itself in good repair in response to injury (Conner et al. 2003).

Several studies have investigated structural modifications of neocortical pyramidal neurons following selective CBF lesions. Injections of 192 IgG-saporin (i.c.v. or i.pr.) induce an increase of apical and basal dendritic spines related to an unaltered dendritic branching in parietal (De Bartolo et al. 2009; Mandolesi et al. 2008) as well as in frontal (Harmon and Wellman 2003; Wellman and Sengelaub 1995) pyramidal neurons during the middle adulthood. The same pattern of changes was found when the selective cholinergic lesion occurred in neonatal age

(De Bartolo et al. 2010). Furthermore, in frontal pyramidal neurons, the lesioninduced increment of dendritic spines has been related to an increased expression of GluR1 subunit of the aminomethylphosphonate (AMPA) receptors (Garret et al. 2006; Kim et al. 2005), suggesting the spines as the morphological substrate for the increase of AMPA receptors induced by the cholinergic lesion. The hypothesis has been advanced that neocortical neurons upregulate AMPA receptors and spine number as a compensatory response to the lesion-induced decreased innervation (Fiala et al. 2002). In fact, the increase in spines would execute a "mechanical" function and prevent lesion-induced atrophy of dendritic branching (Harmon and Wellman 2003). The hypothesis that the spines might provide protection against dendritic atrophy is supported by evidence that in frontal pyramidal neurons of aged rats with forebrain cholinergic lesions, spines did not increase (Harmon and Wellman 2003) and dendritic branching decreased (Works et al. 2004). Nevertheless, an increase in spines does not necessarily imply an increase in functional synapses. A study of slice preparations of hippocampal neurons showed that new spines form even when synapses are inactivated and persist for many hours in the absence of any functional activity (Kirov and Harris 1999). A recent in vivo study (Arellano et al. 2007) reported the existence of non-synaptic spines in the mouse neocortex, indicating that dendritic spines might not be used to form synaptic contacts. A recent review reported that most new spines which appear in adult animals are transient, and only a small percentage of them will ultimately form functional synapses (Alvarez and Sabatini 2007). Moreover, the presence of spatial deficits following forebrain lesions (for details see below) seems to suggest that the newly formed spines of parietal pyramidal neurons are functionally inactive and thus do not directly influence behavior.

The reduced neuroplasticity following CBF-selective lesions is confirmed by many studies investigating the effects of 192 IgG-saporin injections on adult neurogenesis. It has been demonstrated that, 5 weeks after lesion, neurogenesis declined significantly in the granule cell layers of dentate gyrus in hippocampal formation as well as in olfactory bulb. Furthermore, immunotoxic lesions led to increased numbers of apoptotic cells specifically in the subgranular zone, the progenitor regions of dentate gyrus, and within the periglomerular layer of olfactory bulb. So, it has been advanced that the cholinergic system plays a survival-promoting role for neuronal progenitors and immature neurons within regions of adult neurogenesis, similar to effects observed during brain development (Cooper-Kuhn et al. 2004). Moreover, it has been demonstrated that i.c.v. 192 IgG-saporin injections on pnd 7 accelerate the death of newborn cells in the dentate gyrus, but does not change their overall survival rate or phenotypic differentiation (Rennie et al. 2011).

Many effects of selective CBF lesions confirming the reduced neuroplasticity have been described about metabolic markers and are consistent with the view of cholinergic influence over metabolism and amyloid precursor protein (APP) processing in the neocortex and hippocampus. For example, it has been demonstrated that i.c.v. 192 IgG-saporin injections during adulthood causes sustained reductions in glucose utilization in brain regions impacted by cholinergic synapse loss, including frontal cortical and hippocampal regions, relative to glucose use levels (Browne et al. 2001). Also, neurotrophic factors strongly implicated in neuroplasticity display alterations following selective cholinergic depletion. In consequence of i.c.v. 192 IgG-saporin injections, NGF levels increased in the frontal cortex and hippocampus at 3 and 7 days and decreased in the serum at 7 days from lesion, suggesting that opposite changes in brain and serum occur at early stages of cholinergic depletion (Gelfo et al. 2011). 192 IgG-saporin i.c.v. injections induce a reduction of neurotrophin brain-derived neurotrophic factor (BDNF) levels in the parietal cortex as well as in the serum at later stages (15 days from lesion) (Angelucci et al. 2011). It appears interesting, since BDNF has an important role in survival, differentiation and functionality of neurons undergoing degeneration in pathological conditions such as cholinergic neurons in AD.

# 7 Behavioral and Cognitive Outcomes of 192 IgG-Saporin Lesions

The effects of selective immunotoxic lesions have been examined in a variety of behavioral paradigms of learning and memory. However, behavioral studies after 192 IgG-saporin lesions provide a quite unexpected picture: massive cholinergic depletion results in little or no impairment in commonly used spatial and not spatial learning and memory tests (Baxter and Chiba 1999; Everitt and Robbins 1997; McGaughy et al. 2000). Differently, deficits have been noted after cholinergic depletion on tasks that focus on attentional and discriminative abilities (Burk et al. 2002; McGaughy et al. 2002; Ricceri 2003). This general framework is further complicated by route of injection (i.c.v. or i.pr.), lesion extent, age of lesioning, and kind of behavior analyzed.

#### 7.1 Mnesic Functions

The i.c.v. injection method has most consistently produced learning and memory deficits, and the most informative studies have used a range of immunotoxin doses which has allowed analyzing correlations between lesion extent and behavioral deficit.

High-dose (4–10 µg) i.c.v. injections of 192 IgG-saporin result in impairment in the acquisition of navigational learning in Morris Water Maze (MWM) and retention memory deficits in passive avoidance (PA) (Berger-Sweeney et al. 1994; Garcia-Alloza et al. 2006; Leanza et al. 1996; Nilsson et al. 1992; Steckler et al. 1995; Traissard et al. 2007; Waite & Thal 1995; Waite et al. 1995, 1999; Walsh et al. 1995; Zhang et al. 1996). Impaired performance in cued versions of MWM task has been also reported (Berger-Sweeney et al. 1994; Walsh et al. 1995). The behavioral changes are paralleled by a greater than 90 % reduction in CBF neurons that result in 80–90 % reduction in ChAT in the hippocampal and cortical target fields. Conversely, rats with less-extensive CBF lesions following i.c.v. ( $\approx$ 55–75 % cholinergic loss) (Cutuli et al. 2009; De Bartolo et al. 2010; Lehmann et al. 2000, 2002; Mandolesi et al. 2008; Nilsson et al. 1988; Waite et al. 1999) or i.pr. (MS, DBB and/or NBM) immunotoxic injections ( $\approx 15-78$  % cholinergic loss) were only slightly or not at all behaviorally affected in tasks, as MWM, PA, active avoidance, fear conditioning, and T-maze (Baxter et al. 1995, 1996; Dornan et al. 1996; Frick et al. 2004; Garcia-Alloza et al. 2006; Mandolesi et al. 2008; Torres et al. 1994; Wenk et al. 1994).

192 IgG-saporin i.c.v. lesions also impair spatial learning and working memory. A significant correlation between working memory impairment and lesion extent was observed (Wrenn and Wiley 1998; Wrenn et al. 1999), and this dose-dependent impairment was present only in rats that harbored total CBF lesions greater than 75 % (threshold effect). Several studies reported deficient performances in radial arm maze (RAM) (Cutuli et al. 2009; De Bartolo et al. 2010; Lehmann et al. 2003; Traissard et al. 2007) and in a water version of RAM (Antonini et al. 2009) following 192 IgG-saporin i.c.v. injections. Working memory deficits observed in RAM following CBF depletions are consistent with findings obtained by using T-maze alternation test and delayed matching to position/sample (DMTP/DMTS) and delayed nonmatching to position (DNMTP) tasks (Leanza et al. 1996; Lehmann et al. 2000, 2002; McDonald et al. 1997; Paban et al. 2005a, b; Steckler et al. 1995; Wrenn and Wiley 2001). Once again, i.pr. injections producing small, circumscribed cholinergic depletion (<75 %) did not evoke dramatic working memory deficits (McMahan et al. 1997; Pang et al. 2001) or effect in DNMTP task (Steckler et al. 1995). Galani et al. (2002) observed minor impairments only during initial phases of MWM and RAM tests after NBM injection. Perry et al. (2001) found RAM and MWM long-term (11 months) deficits after NBM and MS 192 IgG-saporin lesion, but not short-term deficits (5 months). The few studies considering perseverative behaviors and procedural strategies put into action during RAM demonstrated pervasive deficits of spatial exploration and cognitive planning following i.c.v. (Cutuli et al. 2009; De Bartolo et al. 2010) and NBM (Mandolesi et al. 2008) 192 IgG-saporin lesions.

The role of damage of cerebellar Purkinje cells has been addressed by including additional control groups with selective cerebellar lesions produced by means of OX7-saporin. Injections of 192 IgG-saporin or OX7-saporin (i.c.v.) result in sensory-motor deficits, compatible with the role of Purkinje cells in hyperactivity, motor coordination, and balance (Gandhi et al. 2000; Waite et al. 1995, 1999). Waite et al. (1999) demonstrated that OX7-saporin lesions mimic the effects on MWM acquisition shown after 192 IgG-saporin i.c.v. injections. Reference memory deficits selectively induced by OX7-saporin injections were also found in MWM and RAM (Gandhi et al. 2000; Traissard et al. 2007), confirming the critical importance of cerebellar damage in the i.c.v. lesioning effects. Deficits in the MWM-cued version are addressed to a non-cholinergic cerebellar Purkinje cell damage (Berger-Sweeney et al. 1994; Walsh et al. 1995). On the contrary, Purkinje cell loss is not responsible for PA retention memory and RAM working memory deficits seen after 192 IgGsaporin i.c.v. injections (Wiley et al. 1995; Wrenn et al. 1999). Namely, a threshold relationship between CBF lesion extent and working memory consequences has been demonstrated, indicating that moderate (about 32 %) Purkinje cell loss alone is insufficient to impair RAM performances (Wrenn and Wiley 2001).

As final notation, whether or not a behavioral deficit is observed depends upon an interplay between lesion extent and task demands. For example, recalling a single platform location in the MWM is less demanding than recalling a list of locations in the RAM. Consequently, when lesion extent was high and tasks were demanding, not all 192 IgG-saporin i.pr. studies have yielded negative results in learning and memory paradigms. In fact, Torres et al. (1994) and Rispoli et al. (2006) found impaired PA retention after injection into the NBM. Working memory deficits were reported in MWM (Baxter et al. 1995) and RAM after injections into the MS and/or NBM (Dornan et al. 1996; Mandolesi et al. 2008; Shen et al. 1996; Walsh et al. 1996). Impaired DNMTP performances were found after injection into the MS and the cingulate cortex (Torres et al. 1994; Walsh et al. 1996). Selective removal of the cholinergic projections to the cortical mantle by NBM injections of 192 IgG-saporin led to impaired social interaction reducing the duration of active social contact (Savage et al. 2011).

## 7.2 Spatial/Object Novelty Discrimination

Many studies assessed the effects of 192 IgG-saporin lesions on spatial/object recognition. Unlike the MWM, analysis of the open field (OF) with object performances revealed that i.c.v. 192 IgG-saporin lesions significantly affect spatial discrimination, since lesioned rats did not show selective renewed interest in the displaced objects (Cutuli et al. 2009; De Bartolo et al. 2010; Paban et al. 2005a, b). Conversely, after 192 IgG-saporin i.c.v. and i.pr. injections, the lesioned rats detected the presence of a novel object (Cutuli et al. 2009; De Bartolo et al. 2010; Savage et al. 2011; Vnek et al. 1996). A severe object recognition deficit in rats was observed only after injecting 192 IgG-saporin directly into perirhinal cortex (Winters and Bussey 2005).

The lesion-induced absence of spatial change detection is apparently inconsistent with the tuned MWM performances displayed by the 192 IgG-saporin-lesioned animals. It is possible that the MWM task is more affected by stress levels associated with the forced swimming to search for the escape platform (Cutuli et al. 2009; De Bartolo et al. 2010; Ricceri 2003). The stress responses induced by immersion in water might trigger neuromodulatory compensation that masks mild spatial working memory deficits (Everitt and Robbins 1997; McMahan et al. 1997). In fact, tasks involving high levels of stress may lead to stress-related increases of ACh release in the hippocampus (Gilad 1987; Mizoguchi et al. 2001; Mizuno and Kimura 1997). This could compensate for the greater loss of cholinergic functions in relatively stressful (MWM) than in less stressful (spatial/object novelty) tasks.

#### 7.3 Attentional Functions

Many attentional functions have been found to be sensitive to selective cholinergic depletion, and the robustness of these effects varies with the kind of attentional

demands (McGaughy et al. 2000). Tasks requiring decremental attentional processing, such as latent inhibition, have been found to be unaffected by i.pr. cholinergic depletions (Chiba et al. 1995; Doughtery et al. 1996). However, following NBM or posterior parietal cortex 192 IgG-saporin infusions, subjects were impaired in serial conditioning tasks (Bucci et al. 1998; Chiba et al. 1995). 192 IgG-saporin i.c.v.-lesioned rats were impaired in a 5-Choice Serial Reaction Time test with increasing variability of the delay and visual search demands (Waite et al. 1999). Also the cognitive flexibility consisting in efficiently switching between spatially changing response rules was affected by 192 IgG-saporin i.c.v. lesions, as demonstrated by Cutuli et al. (2009) by using a serial learning task in which rats had to reach a food reward, finding out a sequence of correct open doors daily changed.

The extent of neocortical cholinergic depletion steadily correlates with performance in a sustained attention test (McGaughy et al. 1996, 1999; McGaughy and Sarter 1998, 1999). NBM infusions of 192 IgG-saporin produced dramatic, persistent impairment in the ability to detect unpredictably occurring signals, but not in their correct identification of non-signals in a sustained attentional task. The ability to discriminate signals from non-signals early declined with the additional cognitive demand provided by prolonged time on task. Dose-dependent cholinergic cortical denervation after NMB or i.c.v. lesions resulted in consequent behavioral impairments. Similar sensitivity to cortical cholinergic denervation was found when divided attention was tested. In conditional discrimination tests, lesioned animals showed increased response latencies in conditions requiring the simultaneous maintenance of two sets of conditional response rules (Turchi and Sarter 1997; Waite et al. 1999).

# 7.4 Neonatal Lesions

Selective cholinergic lesions by means of 192 IgG-saporin have been performed also in developing rats. Neonatal 192 IgG-saporin i.c.v. lesions produce a permanent loss in CBF, without inducing the cerebellar damage shown following i.c.v. infusion in adults.

While on the first postnatal day (pnd 1) single 192 IgG-saporin i.c.v. injections induce a very mild cholinergic loss in the cortex (17 % reduction in cortical ChAT activity) accompanied by no reduction in hippocampal ChAT activity and no behavioral alteration (Ricceri 2003; Ricceri et al. 1997, 1999), combined pnd 1 and 3 192 IgG-saporin injections lead to a dramatic reduction of ascending cholinergic projections to both the hippocampus and neocortex and impaired PA learning and MWM (Leanza et al. 1996; Ricceri et al. 2002). After 192 IgG-saporin i.c.v. injections at pnd 4, ChAT activity was markedly reduced in the hippocampus and cortical regions (73–84 %), but rats were unimpaired in MWM (Leanza et al. 1996). At pnd 7 192 IgG-saporin injections provoked consistent reductions in ChAT activity (78–84 % hippocampus, 52–64 % neocortex) accompanied by early and long-term behavioral impairments (Ricceri 2003; Wrenn and Wiley 1998).

In fact, pnd 7-lesioned rats displayed altered ultrasonic vocalizations in maternal separation and fear conditioning paradigms (Ricceri 2003; Ricceri et al. 2007; Scattoni et al. 2005), learning deficits in PA, and reduced exploratory behavior (Ricceri et al. 1997). At adulthood pnd 7-lesioned rats showed no deficits in MWM and delayed spatial alternation T-maze (De Bartolo et al. 2010; Pappas et al. 1996; Ricceri et al. 1999) but altered reactions to spatial novelty (De Bartolo et al. 2010; Ricceri et al. 1999), deficits in social transmission of food preferences (Ricceri et al. 2004), impaired working memory performances in RAM, and impulsive behaviors (De Bartolo et al. 2010; Pappas et al. 2005; Scattoni et al. 2006). In summary, as in adult animals, neonatal 192 IgG-saporin i.c.v. injections result in behavioral impairment only when the resultant CBF lesion is extensive and tasks are highly demanding.

## 8 Conclusion

The immunotoxin 192 IgG-saporin causes a selective, long-lasting loss of the cholinergic neurons within the CBF and is able to interfere with experiencedependent plasticity. A number of studies have demonstrated alterations of several structural and biochemical parameters related with neuroplasticity (dendritic spines and branching of pyramidal neurons, adult neurogenesis, levels of neurotrophic factors) in both cortical mantle and hippocampus.

The introduction of immunotoxic lesioning methods has proved to be a valuable tool in the investigation of the role of ascending cholinergic systems in cognition. The effects of selective immunotoxic lesions have been examined in a variety of behavioral paradigms. The general framework has to take into account the route of injection (i.c.v. or i.pr.), lesion extent, age of lesioning, and kind of behavior analyzed in the different studies. Overall, cholinergic depletion has been demonstrated to provoke selective learning and memory impairments as well as deficits in attentional and discriminative abilities. However, as demonstrated by the literature reviewed in this chapter, the specificity of the neurotoxic lesioning method is strictly linked to the validity and sensitivity of behavioral methods. Furthermore, 192 IgG-saporin lesions result in overt behavioral deficits only with high-grade CBF lesions, indicating a threshold relationship between CBF lesion extent and cognitive impairment.

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# TRPV1 Activators ("Vanilloids") as Neurotoxins

# Ashutosh Kumar, Rakesh Kumar Majhi, Manoj Yadav, Arpad Szallasi, and Chandan Goswami

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#### Abstract

A distinct subset of primary sensory neurons is distinguished by their unique sensitivity to capsaicin, the pungent ingredient in hot chili peppers. The initial excitation by capsaicin of these neurons is followed by a long-lasting, but fully reversible, refractory state (traditionally termed as desensitization) or under certain conditions, like neonatal treatment, frank neurotoxicity. This neurotoxic action was extensively used to identify capsaicin-sensitive neuronal pathways and to explore their physiological function. In 1997, a specific receptor for capsaicin and related compounds (collectively referred to as vanilloids) was identified as transient receptor potential cation

A. Kumar • R.K. Majhi • M. Yadav • C. Goswami (🖂)

e-mail: chandan@niser.ac.in

Authors Ashutosh Kumar and Rakesh Kumar Majhi have equally contributed.

National Institute of Science Education and Research, Institute of Physics Campus, Bhubaneswar, Orissa, India

A. Szallasi (⊠) Monmouth Medical Center, Long Branch, NJ, USA e-mail: ASzallasi@barnabashealth.org; ASzallasi@sbhcs.com

channel subfamily V member 1 (TRPV1), a multifunctional channel involved in thermosensation (heat) and taste perception (e.g., peppers and vinegar). Importantly, TRPV1 also functions as a molecular integrator for a broad range of seemingly unrelated noxious stimuli including venoms from spiders and jellyfish. Indeed, TRPV1 is thought to be a major transducer of the thermal hyperalgesia that follows inflammation and/or tissue injury. Ablation of sensory neurons by vanilloids is, however, not only a research tool but also has a clear therapeutic potential. Currently, sitespecific resiniferatoxin (an ultrapotent capsaicin analog) injections are being evaluated as "molecular scalpels" to achieve permanent analgesia in cancer patients with chronic, intractable pain. In this chapter, we review our knowledge of the molecular mechanisms underlying vanilloid-induced neurotoxicity, which includes both TRPV1-mediated and independent signalling pathways.

#### Keywords

Capsaicin • Resiniferatoxin • The capsaicin (vanilloid) receptor TRPV1 • Vanilloids

List of Abbrevia	tions
CCK-B	Cholecystokinin receptor-B
CGRP	Calcitonin gene-related peptide
CNS	Central nervous system
CTB	Cholera toxin subunit B
DRG neuron	Dorsal root ganglion neuron
ETC1	Electron transport chain complex 1
ETC3	Electron transport chain complex 3
ICK-peptides	Inhibitory cystine knot peptides
ISH	In situ hybridization
MPT	Membrane permeability transition
NADA	N-arachidonoyl-dopamine
NNOS	Neuronal nitric oxide synthase
OLDA	N-oleoyldopamine
PALDA	N-palmitoyldopamine
PCR	Polymerase chain reaction
РКС	Protein kinase C
PMOR	Plasma membrane NADH oxidoreductase
ROPA	Resiniferonol 9,13,14-orthophenylacetate
ROS	Reactive oxygen species
RTX	Resiniferatoxin
SP	Substance P
STEARDA	N-stearoyldopamine
TG-neurons	Trigeminal ganglion neurons

TiTX	Tinyatoxin	1				
TRPV1	Transient	receptor	potential	cation	channel	subfamily
	V member	1				

#### 1 Introduction

Natural products provide a window of opportunity to identify new targets for pharmacological intervention. Capsaicin, the active principle in hot chili peppers (*Capsicum annum*), is a prime example. Connoisseurs of hot spicy food are intimately familiar with the predominant pharmacological actions of capsaicin from personal experience: it induces profuse perspiration (known as gustatory sweating) as well as a hot, burning sensation that dissipates upon repeated challenge (desensitization). Evolutionary selective pressure seems to have maximized the pungency of capsaicin. It was speculated that the compound's pungency is able to deter ambulatory animals from eating chili pepper fruits, favoring those plants whose seeds were dispersed widely by birds. Indeed, the avian TRPV1 receptor is not activated by capsaicin, and hence birds are undeterred from ingesting chili pepper fruits and can excrete the pepper seeds large distances away. This forms the basis of the development of hot pepper-flavored "squirrel-free" bird feed. It is still a mystery, however, why the same pungency that repels squirrels is perceived as pleasurable by many human beings.

Although topical capsaicin has been used in folk medicine to relieve pain for centuries, it took the genius of the late Miklós (Nicholas) Jancsó to ask the obvious question: how can an irritant molecule paradoxically cause analgesia? Jancsó noticed that the initial transient irritation by capsaicin was followed by a long-lasting refractory state in which the neurons not only did not respond to a repeated capsaicin challenge but were also insensitive to various unrelated stimuli, and he termed this state as "desensitization" (Jancsó and Jancsó 1949; Jancsó-Gábor et al. 1970). Per definition, capsaicin-mediated desensitization is reversible. However, following neonatal treatment or when sufficiently high doses are used in adult animals, capsaicin induces neuronal death and causes permanent analgesia (Jancsó et al. 1984). This forms the experimental foundation for the use of site-specific resiniferatoxin (an ultrapotent capsaicin analog) injections to achieve permanent relief of otherwise intractable pain in cancer patients. Furthermore, chemical ablation of sensitive neurons by capsaicin turned out to be an invaluable tool to dissect capsaicin-sensitive pathways and to explore their physiological function. Generally speaking, these neurons comprise a functional subset of peptidergic primary sensory neurons with unmyelinated (C-type) fibers that respond to noxious stimuli and initiate the cascade of neurogenic inflammation.

The molecular mechanisms underlying capsaicin-mediated desensitization are poorly understood, and the line between desensitization and neurotoxicity is most likely ill defined and arbitrary. The sensitization of TRPV1 by different kinases and involvement of different pathways also remain unclear (Vellani et al. 2006; Otten et al. 1983; Negri et al. 2006). The cloned capsaicin (vanilloid) receptor TRPV1 is an inwardly rectifying cation channel with limited selectivity for  $Ca^{2+}$ , and  $Ca^{2+}$ -overload is a well-established cause of neurotoxicity (Chard et al. 1995; Caterina et al. 1997; Sugimoto et al. 1998; Czaja et al. 2008). Indeed, neurochemical calcium and other ions such as cobalt stains were historically used to visualize sensitive neurons in capsaicin-treated animals (Winter 1987). At the ultrastructural level, neuronal calcium staining is coupled to swollen mitochondria, but it is still hotly debated if this reflects a buffering of cytoplasmic  $Ca^{2+}$  by the mitochondria (a direct TRPV1-mediated effect) or an indirect, TRPV1-independent interference of capsaicin with the mitochondrial electron transport chains (ETCs) (Szolcsányi et al. 1971).

The first vanilloids to be identified were pungent phytochemical compounds that typically contain a vanillyl (4-hydroxy-3-methoxybenzyl) moiety essential for bioactivity (hence the name "vanilloid") and are present in nature, mainly as plant resources (Table 1). Prime examples include capsaicin (the pungent ingredient in chili peppers), piperine (responsible for the piquancy of black pepper), resiniferatoxin (isolated from the latex of the cactus-like plant Euphorbia resinifera), curcumin (from Curcuma longa), eugenol (in clove), and zingerone (from ginger). Subsequently, vanilloids (so-called endovanilloids) were isolated from animal tissues (e.g., N-arachidonoyl-dopamine, NADA, isolated from rat brain) or were obtained via chemical synthesis to explore structure-activity relations (Huang et al. 2002). Few more examples are N-oleoyldopamine (OLDA). *N*-palmitoyldopamine (PALDA), and *N*-stearoyldopamine (STEARDA) (Chu et al. 2003). Indeed, the existence of a specific capsaicin receptor was first postulated based on the fairly strict structural requirements for capsaicin-like bioactivity.

# 2 Capsaicin-Sensitive Neurons

In culture, capsaicin kills ~35 % of dorsal root ganglion (DRG) neurons. It was postulated that capsaicin destroys these neurons due to influx of excess  $Ca^{2+}$ (Chard et al. 1995). Indeed, capsaicin causes a drastic increase in basal intracellular  $Ca^{2+}$  concentration when applied to the cultured DRG neuron. This in vitro toxicity correlates with the observation that application of capsaicin in vivo for prolonged periods can cause irreversible toxic effect leading to loss of painsensing neurons (Holzer 1991; Maggi 1991). The DRG neurons abolished by capsaicin are small in size, possess either unmyelinated C- or thinly myelinated  $A\delta$ -fibers, and show large inward intracellular  $Ca^{2+}$  currents in response to capsaicin. Both  $Ca^{2+}$  current and the neurotoxic effect of capsaicin were prevented by pre-treating the cells with ruthenium red, currently recognized as a general, nonspecific blocker of TRP channels. The neurotoxic action of capsaicin can also be prevented by chelating extracellular  $Ca^{2+}$  and/or by blocking  $Ca^{2+}$ activated proteases like calcineurin (Docherty et al. 1996). Taken together, these results imply that capsaicin exerts its neurotoxic effects predominantly due to

Table 1 Exam	Table 1 Example of some vanilloids				
Vanilloids	Chemical structure	Molecular formula	Origin	Properties and action	Reference
Capsaicin	H H O H	C <sub>18</sub> H <sub>27</sub> NO <sub>3</sub> Capsicum genus	<i>Capsicum</i> genus	Analgesic, anticancer, antiobesity, anti-inflammatory, antioxidant activates TRPV1 at 10-8 to 10-6 M	Caterina et al. 1997; Luo et al. 2011; Knotkova et al. 2008.
Resiniferatoxin		C <sub>37</sub> H <sub>40</sub> O <sub>9</sub>	Euphorbia resinifera	Inhibits inflammatory hyperalgesia, activates TRPV1 at 10-10 to 10-8 M	Neubert et al. 2008; Caterina et al. 1997.
Paradol		C <sub>17</sub> H <sub>26</sub> O <sub>3</sub>	Aframomum melegueta	Antioxidative, antitumor, activates TRPA1 and TRPV1	Chung et al. 2001; Riera et al. 2009.
					(continued)

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Vanilloids	Chemical structure	Molecular formula	Origin	Properties and action	Reference
Curcumin	e e e e e e e e e e e e e e	$C_{21}H_{20}O_{6}$	Zingiberaceae family	<i>Zingiberaceae</i> Used for pain relief, anticarcinogenic family and food additive	Kuptniratsaikul et al. 2000; Aggarwal and Shishodia 2004.
Rotenone	H <sup>3</sup> COCH <sup>3</sup> H <sup>3</sup> COCH <sup>3</sup> H <sup>3</sup> COCH <sup>3</sup>	C <sub>23</sub> H <sub>22</sub> O <sub>6</sub>	Lonchocarpus nicou	C <sub>23</sub> H <sub>22</sub> O <sub>6</sub> Lonchocarpus Used as pesticide, insecticide, <i>nicou</i> piscicide, and neurotoxin	Swarnkar et al. 2012; Cabeza-Arvelaiz and Schiestl 2012
Coenzyme Q	H <sub>3</sub> CO H <sub>3</sub> CO CH <sub>3</sub> CH <sub>3</sub>	C <sub>59</sub> H <sub>90</sub> O <sub>4</sub> Most eukar cells	Most eukaryotic cells	Antioxidant	Littarru and Tiano 2007; Hoppe et al. 1999

consequences of both increased  $Ca^{2+}$  concentration inside the cell and cascades of cellular signalling events (Chard et al. 1995).

Capsaicin-sensitive sensory neurons are bipolar neurons with somata in dorsal root and trigeminal ganglia. The peripheral fibers innervate the skin as well as the viscera, whereas the central axons enter the central nervous system (CNS) where they form synapse at second-order neurons in the dorsal horn of the spinal cord and the spinal nucleus of the trigeminal tract, respectively. The vagal nerve also supplies capsaicin-sensitive innervations to some visceral organs: the cell bodies of these neurons are in the nodose ganglia, and they centrally project to the area postrema. The presence of TRPV1 in these neuronal pathways is firmly established by a combination of [<sup>3</sup>H]-labeled resiniferatoxin (RTX) autoradiography, TRPV1 immunostaining, and other molecular methods (Szallasi and Blumberg 1999). The existence of TRPV1-expressing brain nuclei is, however, controversial.

In 1988, using a silver impregnation method, Ritter and Dinh described unexpected neuronal degeneration in several discrete forebrain and hindbrain areas of adult rats following i.p. capsaicin administration (50 or 90 mg/kg) (Ritter and Dinh. 1988). Subsequently, the presence of TRPV1 receptors was reported in corresponding (and also in additional) rat brain nuclei (and also in other parts) and the cortex in  $[^{3}H]RTX$ -binding experiments as well as by polymerase chain reaction (PCR), in situ hybridization (ISH), Western blot analysis, and immunostaining studies targeting to detect TRPV1 at RNA and protein level (Acs et al. 1996; Tóth et al. 2005; Doyle et al. 2002; Valtschanoff et al. 2001; Caterina et al. 1997; Goswami et al. 2010; Sasamura et al. 1998). Recordings obtained from rat brain slices implied that these TRPV1 receptors were functional and might be involved in learning and memory formation. Indeed, TRPV1 has been linked to the pathogenesis of Alzheimer disease (Micale et al. 2010; Pákáski et al. 2009). An additional important role for brain TRPV1 in anxiety and fear was postulated based on studies in TRPV1 knockout animals (Marsch et al. 2007). Most recently, however, only weak and minimal TRPV1 expression was found in rodent brain using a sensitive and selective reporter-mouse model (Tóth et al. 2005). This discrepancy is puzzling. Furthermore, relevant to the topic of this review, it clearly questions the value of capsaicin-induced toxicity to dissect "capsaicin sensitive" (i.e., TRPV1-expressing) pathways.

#### 3 Plant-Derived Vanilloid Toxins

Though there are handful examples of vanilloids from plant origin, in this chapter, only vanilloids, namely, capsinoid and resiniferanoids, will be discussed in details.

*Capsinoids*: Capsaicin, the archetypal vanilloid, is responsible for the piquancy of hot chili peppers (Buck and Burks 1986). Capsaicin and related compounds are collectively called capsinoids. Naturally occurring capsinoids include capsiate, dihydrocapsiate, and nordihydrocapsiate. Like capsaicin, capsinoids activate TRPV1 (Iida et al. 2003), yet, they were reported to be sweet-tasting. Of synthetic capsinoids,

in the past olvanil and nuvanil had attracted attention as "improved" (candidate molecules that can be given per os to achieve desensitization) in preclinical models of chronic pain. The synthetic capsaicin, civamide (*cis*-capsaicin), is under clinical development by Winston Pharmaceuticals for indications like cluster headache.

Resiniferanoids: Resiniferatoxin (RTX) is a naturally occurring ultrapotent capsaicin analog isolated from the dried latex of the cactus-like perennial Euphorbia resinifera, a native of the Anti-Atlas Mountains in Morocco (Appendino and Szallasi 1997). The closely related tinyatoxin (TiTX) is present in the soap of Euphorbia poissonii, a succulent plant in Nigeria, which is traditionally used as a pesticide. TiTX is somewhat (approximately threefold) less potent than RTX. Although a method for the full synthesis of RTX is available, the molecule is still isolated from its natural host or obtained semi-synthetically from its parent diterpene, resiniferonol 9,13,14-orthophenylacetate (ROPA). Attempts to recapitulate the ultra potency of RTX by synthesizing phorbol-based vanilloids (phorbol is inexpensive and its chemistry is well known) met limited success. Of note, ROPA is a potent activator of protein kinase C (PKC) and acts as a tumor promoter (Frey et al. 2004). At nanomolar concentrations RTX activates TRPV1, but does not activate PKC, and does not promote the formation of tumor either in the mouse skin model of two-stage carcinogenesis (Driedger and Blumberg 1980). RTX, however, did show unexpected activity against various cancer cell lines in a most likely TRPV1-independent manner (discussed later in details).

## 4 Venoms as Toxins for Vanilloid Receptors

Venomous creatures such as spiders, scorpions, snakes, and snails contain a wealth of peptide toxins which can activate or block different ion channels including vanilloid receptors and produce shock, paralysis, or death (Escoubas and Rash 2004; Miller 1995; Terlau and Olivera 2004). Often such venoms and toxins and their activity on ion channels define the pray-predator relationship and exert selection pressure of certain species in specific environmental niches (Fry et al. 2006). Some venoms contain toxins which act on somatosensory neurons and provoke a noxious sensation, primarily in mammals (Siemens et al. 2006; Bohlen and Julius 2012). Somewhat unexpectedly, bites or stings by such venomous creatures were shown to activate the TRPV1 receptor and produce inflammatory pain (Siemens et al. 2006; Cuypers et al. 2006).

Venom from a *Tarantula* sp., *Psalmopoeus cambridgei*, commonly found in the West Indies, has three inhibitory cystine knot (ICK) peptides present in one peptide (Siemens et al. 2006). Previously, it was shown that venoms and toxins from spiders, snakes, and scorpions mainly target three members of the TRP channel family, namely, TRPV1, TRPA1, and TRPM8 (Siemens et al. 2006). By reverse phase chromatographic technique, the *Tarantula* venom was purified and fraction-ated further. When testing the bioactive fractions on HEK-293 cells heterogeneously expressing different TRP channels, it was demonstrated that the toxin was selective to TRPV1 (Siemens et al. 2006). Edman sequencing revealed the

presence of three closely related peptides named vanillotoxin 1, vanillotoxin 2, and vanillotoxin 3, all of which can act as agonists for TRPV1. Similar to the ICK peptide, these vanillotoxins have six cysteine residues in each toxin sequence (Zhu et al. 2003). Recently, another tarantula, namely, *Ornithoctonus huwena*, from China has been reported to produce a toxin that functions as a TRPV1 agonist (Bohlen et al. 2010). Interestingly, this toxin is similar to the vanillotoxin 1–3. This toxin is named double knot toxin because it has two ISK motifs repeated in tandem (Bohlen et al. 2010).

For the sake of completeness, it should be mentioned that there are venoms which contain inhibitory components against TRPV1. For example, extract from a nematocyte (sea anemone *Heteractis crispa*) contains a peptide, named APHC1, which blocks TRPV1 (Andreev et al. 2008). Furthermore, venom from a funnel web spider (*Agelenopsis aperta*) has two toxin components (AG489 and AG505) which inhibit TRPV1 (Kitaguchi and Swartz 2005).

## 5 The Diversity of Vanilloid Actions

The spectrum of vanilloid actions depends not only on the pharmacophore but also on the species and the route of administration. One has to carefully consider all these factors before selecting the appropriate compound for the studies.

Pharmacophore: RTX functions as an ultrapotent capsaicin analog with a peculiar spectrum of pharmacological activities. For example, in the rat, RTX is several thousandfold more potent than capsaicin in inhibiting the xylene-induced neurogenic inflammatory response or in causing twitch inhibition in the vas deferens (Wardle et al. 1996). By contrast, RTX is only similar in potency to capsaicin in evoking the eye-wiping response upon intraocular instillation. RTX also has unique actions: unlike capsaicin which repeatedly evokes the pulmonary chemo-reflex with no evidence of desensitization, RTX desensitizes this reflex without causing any detectable prior activation. Interestingly, RTX and capsaicin congeners showed distinct structure-activity relations for receptor binding and Ca<sup>2+</sup> uptake, respectively. Resiniferanoids displayed high affinity in the binding assay and low activity in the  $Ca^{2+}$  uptake assay, whereas capsinoids showed the opposite pattern. Based on these observations, the existence of two types of vanilloid receptor was postulated, an R-type (presumably metabotropic) vanilloid receptor responsible for desensitization and a C-type (most likely) ionotropic receptor mediating irritation. This hypothesis, however, turned out to be not true: following the cloning of TRPV1, it was recognized that the same receptor mediates both R-type binding and C-type calcium responses (Caterina et al. 1997). Pharmacokinetics appears to play an important role in determining RTX-like and capsaicin-like activities. In most cases, capsaicin-evoked Ca<sup>2+</sup> current is rapid in onset, large in magnitude, and short in duration. By contrast, RTX induces sustained, slowly developing and long-lasting currents. This observation might explain why RTX treatment shows a better desensitization-to-excitation ratio than capsaicin for most end-points examined.

Species Differences: Evolutionary, heat, and capsaicin sensitivity seems to have developed over a time in different species. However, different species reveal differences in responsiveness against heat and capsaicin (Nagy and Rang 2000; Phillips et al. 2004; Jordt and Julius 2002; Sardar et al. 2012). Even mammalian species show striking differences in the way they react to capsaicin challenge (Gavva et al. 2004). Reptiles, such as snakes, seem to be insensitive to capsaicin as trigeminal ganglion (TG) neurons from snakes do not respond to capsaicin (Gracheva et al. 2010). Birds do not respond to capsaicin. In keeping with this, chicken DRG neurons lack specific [<sup>3</sup>H]RTX-binding sites, and the treatment of DRG neurons from chicken with capsaicin does not results in  $Ca^{2+}$  influx (Jordt and Julius 2002; Goswami et al. 2007). Chicken, however, do possess TRPV1 but it is resistant to capsaic due to a point mutation at the position of S512 which confers capsaicin sensitivity in mammals (Jordt and Julius 2002). Within mammals, rats and rabbits are a good example for species-related differences in capsaicin sensitivity (Gavva et al. 2004). In rats, application of 1 % capsaicin to the saphenous nerve resulted in a 35 % decrease in the C:Aδ fiber ratio (Pini et al. 1990), while the same concentration of capsaicin administration to saphenous nerve of rabbits did not change the C: A $\delta$  fiber ratio (Lynn and Shakhanbeh 1988). This implies the existence of low density and/or low capsaicin-affinity TRPV1 receptors in the rabbit. Indeed, no high-affinity RTX binding was detected in rabbit DRG neuron membranes (Gavva et al. 2004). An intriguing species-related difference in RTX actions was reported in the cat. Unlike in rats where, as described above, RTX desensitizes the pulmonary chemoreflex without prior activation, in the cat, RTX evokes the full reflex triad (Pórszász and Szolcsányi 1991–1992; Sculptoreanu et al. 2005). Bat has two isoforms of TRPV1 of which one is responsive to infrared (Gracheva et al. 2011). However, both isoforms are responsive to capsaicin and high temperature.

Routes of Administration: The site of administration of vanilloids can be an important factor in determining neurotoxicity. This is due to the fact that intrathecal, perineural, intraganglionic, subcutaneous, intraperitoneal, and topical application of vanilloids exposes different parts of sensory neurons to differential vanilloid concentrations. For example, intraganglionic injections expose the cell bodies of sensitive neurons to high vanilloid concentrations, leaving the peripheral terminals intact. By contrast, topical vanilloid administration mainly targets the peripheral nerve terminals with no detectable effect on perikarya. As an added complication, different parts of the primary afferent neurons (central terminals, cell bodies, axons, and peripheral terminals) have their own range of sensitivity to vanilloid agonists. For example, peripheral sensory axons have different sensitivity to capsaicin, with the most vulnerable part being the subepidermal part of the axon (Chung et al. 1990). Intrathecal or intraganglionic administration of RTX selectively deletes TRPV1-expressing neurons and confers irreversible thermal, inflammatory, and cancer pain relief (Brown et al. 2005; Karai et al. 2004; Tender et al. 2005). By contrast, peripheral administration of a single dose of RTX produces a longlasting but reversible analgesia by ablating nociceptive nerve terminals (Karai et al. 2004; Kissin 2008; Neubert et al. 2003; Goswami et al. 2007). This effect could last from few days to few weeks and is preceded by a rapid loss of specific [<sup>3</sup>H]RTX-binding sites which is already apparent 1 h after RTX treatment. Similarly, a long-lasting (up to 1 month) but fully reversible loss of TRPV1-like immunoreactivity was observed in skin biopsies taken from volunteers following exposure to a high-concentration capsaicin patch (Qutenza) (Inoue et al. 2002). Taken together, these results imply that it is impossible to extrapolate vanilloid actions from one species to another, especially if the route of administration is also changed.

#### 6 Vanilloid Interaction with Receptors

Among all TRPV channels, TRPV1 represents the best characterized channel in terms of the vanilloid-mediated activation. It has been demonstrated that the temperature gating of TRPV1 is actually a voltage-dependent process (Voets et al. 2005). In a similar manner, vanilloids also modulate the voltage dependency of the TRP channel in ambient temperature and thus eliminates the requirement for high temperatures. In a simplified manner, it can be said that at molecular level, interaction of vanilloids with TRPV1 results in a conformational change which reduces the requirements for high temperature and thus is suitable for channel opening. However, information regarding the species-specific interaction of vanilloids to TRPV1 remains insufficient and discrepancies in the vanilloid-binding sites remain ambiguous.

Initially, it was demonstrated that capsaicin and capsazepine are competitive inhibitors of RTX binding on membrane fraction suggesting that these compounds probably bind at the same region (Szallasi et al. 1993). Capsaicin was originally believed to activate TRPV1 by binding to its extracellular loops. However, use of a membrane impermeable analog of capsaicin, i.e., DA-5018.HCl, reveals opposite effect on the TRPV1 present in the inside-out patched membrane. This compound activates TRPV1 when applied outside of the pipet, but failed to activate TRPV1 when applied from inside of the patch. This result confirms that capsaicin, being lipophilic in nature, can cross the plasma membrane and thus can bind to an intracellular/intramembranous region of TRPV1 (Jung et al. 1999). Indeed, it was demonstrated that capsaicin binds to the TM regions of TRPV1. The capsaicinbinding region was mapped down to a critical residue (Y511) located at the region between the 1st intracellular loop joining TM2 and TM3 regions of rat TRPV1 (Jordt and Julius 2002). The same position is responsible for capsaicin sensitivity too. In agreement with the involvement of that residues in capsaicin sensitivity, two mutants, namely, TRPV1-Y511A and TRPV1-S512Y (rat TRPV1), abolished capsaicin-mediated activation, while TRPV1-Y511A mutant failed to respond to anandamide also (an endogenous vanilloid that activates TRPV1) (Jordt and Julius 2002). Functional studies with the capsaicin-insensitive orthologs of TRPV1 (avian and rabbit) have demonstrated that capsaicin-mediated gating requires a small region of the TRPV1 which includes residues at positions 511 and 512 (Jordt and Julius 2002; Gavva et al. 2004). Interestingly, the mutant, namely, TRPV1-S512Y, converted the activity of the antagonist 5'I-RTX into an intrinsic agonist, albeit with a lower potency than its parent compound, RTX (Sutton et al. 2005). Notably, these two residues are present between transmembrane domain 3 (TM3) and the first intracellular loop region. Based on these observations, it was proposed that gating of this channel may involve a sequential movement of a paddle structure. According to this concept, the TM3 and TM4 region of the channel is predicted to form a gating paddle (Chou et al. 2004), with residues such as M547 (Rat) and L547 (human) forming a part of the key agonist-binding site which is accessible from the intracellular interface (Johnson et al. 2006). However, this suggests that the critical agonist-binding region is not buried deep within the transmembrane region as suggested by the traditional homology models (Jordt and Julius 2002; Gavva et al. 2004; Phillips et al. 2004). This concept is also supported by the recent observation made in voltage-dependent potassium channel KvAP where X-ray crystal structure confirmed that amino acids at position 512 and 547 are located in close proximity (Jiang et al. 2003). In agreement with the joint action of TM3 and TM4 in agonist recognition, residue M/L547 located at the TM4 mediates significant species differences in resiniferatoxin (RTX) sensitivity, and the S512 is critical for discriminating between pH and capsaicin gating of TRPV1 (Jordt and Julius 2002).

However, a number of studies indicated the involvement of additional residues located within TM4, the putative S5-S6 pore region, and also at the N- and C-termini of TRPV1 on vanilloid-mediated activities and capsaicin responsiveness (Welch et al. 2000; Vlachova et al. 2003; Gavva et al. 2004; Phillips et al. 2004; Jung et al. 2002). For example, involvement of residues (R114 and E761) located in the N- and C-cytosolic tails of rat TRPV1 respectively has influence on the RTXbinding and RTX-mediated response to TRPV1 (Jung et al. 2002). Similarly, a number of critical residues located within the TM3 and TM4 (also considered as voltage sensor) regions have been shown to be responsible for major speciesspecific differences in vanilloid activity (Chou et al. 2004; Gavva et al. 2004; Phillips et al. 2004). In agreement with the involvement of TM3 and TM4 in the capsaicin activity in different species, M547 and T550 located in TM3 and TM4 region of TRPV1 (rat as well as human) confer vanilloid sensitivity, [<sup>3</sup>H]RTXbinding and capsazepine-binding, parameters which are different in rabbit TRPV1 (Gavva et al. 2004). Changing the single residue at 550 in rabbit TRPV1 to the corresponding residue found in rat and human TRPV1 (I550T) was sufficient to confer gain of function for activation by capsaicin (Gavva et al. 2004). Furthermore, TRPV1 (rat as well as in human) mutants, namely, TRPV1-T550I and TRPV1-Y511A, demonstrate a loss of sensitivity to capsaicin (Gavva et al. 2004). A single mutation embedded in the TM4 region of TRPV1 (human), namely, TRPV1-L547M, produced a 30-fold increase in sensitivity to [<sup>3</sup>H]RTX, whereas the reverse mutation in the rat isoform caused a decrease in sensitivity of equal amplitude (Chou et al. 2004). These critical residues affect ligand recognition to some extent and also affect channel function in response to ligand binding (Gavva et al. 2004).

Three amino acid residues located near the pore region of the rat TRPV1 are also involved in the capsaicin responsiveness. This is confirmed by the TRPV1 mutants, namely, TRPV1-E636Q, TRPV1-D646N, and TRPV1-E648Q mutants, which affects capsaicin-mediated gating but not the heat or proton-mediated activation of TRPV1 (Welch et al. 2000). In addition, mutation of three residues located in TM6 (NML676FAP) also abolished capsaicin-mediated activation with little effect on ligand binding to rat TRPV1 (Kuzhikandathil et al. 2001). Similarly, another mutant, namely, TRPV1-L547M (Human) located at the 4th TM region, causes a decrease in capsaicin potency (Johnson et al. 2006).

#### 7 Vanilloid-Induced Messenger Plasticity

It was postulated that vanilloids do not affect non-nociceptive neurons and/or mechanosensitive nociceptive neurons (Karai et al. 2004). This is important because, unlike local anaesthetics which target ubiquitous sodium channels in all axons, vanilloids selectively block heat-sensitive TRPV1-positive nociceptors and thus leave other sensory modalities intact, rendering these molecules a better choice for pain relief. In the rat spinal cord, RTX treatment ameliorated the so-called "wind-up" phenomenon after repeated peripheral C-fiber stimulation that is believed to correspond to central sensitization (Xu et al. 1997). This effect on the "wind-up" response correlated to the time course of the thermal hypoalgesia in the hot plate test (Xu et al. 1997). Unexpectedly, RTX treatment also resulted in reduced sensitivity to mechanical stimulation (Pan et al. 2003). The mechanical hypoalgesia was, however, transient (a few days) compared to the thermal hypoalgesia (several weeks). Since TRPV1-expressing C-fiber neurons do not respond to mechanical stimuli (and mechanosensitive A neurons do not express TRPV1), it is unclear how RTX treatment elevates the threshold for mechanical stimulation. One might wonder, however, if this is somehow part of the "messenger plasticity" that follows vanilloid administration.

Paradoxically, RTX treatment can cause mechanical allodynia. Pan et al. injected  $200 \,\mu g/kg \,RTX$  i.p. to rats and observed rapid increase in the paw withdrawal latency to a heat stimulus, while profound tactile allodynia developed in 3 weeks (Pan et al. 2003). This unexpected increase in mechanical sensitivity lasted for at least 6 weeks (Pan et al. 2003). In the RTX-treated rats, IB4-labeled unmyelinated C-fiber terminals in the dorsal horn were significantly reduced, and cholera toxin subunit B (CTB)labeled myelinated fiber terminals appeared to sprout into lamina II of the spinal dorsal horn (Pan et al. 2003). Electron microscopic examination of the sciatic nerve also revealed significant loss of unmyelinated fibers and extensive ultrastructural damage of myelinated fibers in RTX-treated rats (Pan et al. 2003). Immunofluorescence labeling showed diminished TRPV1-like immunoreactivity in DRG neurons and the spinal dorsal horn following RTX administration. This study suggests that systemic (i.p.) RTX administration diminishes the thermal pain sensitivity by depletion of unmyelinated afferent fibers. At the same time, RTX damages the myelinated afferent fibers and causes their abnormal sprouting in lamina II of the spinal dorsal horn. The latter anatomical change might be the reason behind delayed tactile allodynia (Pan et al. 2003).

As discussed above, desensitization to vanilloids is reversible and thus contrasts to irreversible neurotoxicity. For example, it has been reported that a single topical application of RTX to the rat cornea reduces the capsaicin-evoked eve-wiping response in a dose-dependent manner for 3–5 days, while the normal nociceptive responses return by 5-7 days (Bates et al. 2010). Importantly, RTX administration did not impair epithelial wound healing and blink reflex or cause detectable histological damage to the cornea (Bates et al. 2010). Immunohistochemistry experiments revealed that RTX treatment caused a temporary loss of calcitonin gene-related peptide (CGRP) expressing nociceptive fibers: the majority of fibers reappeared within 12 days and full recovery was attained within 4 months (Bates et al. 2010). Systemic (s.c.) RTX treatment also inhibits the capsaicin-induced eyewiping response and depletes CGRP-like immuoreactivity in the dorsal horn of the spinal cord: these effects last much longer (months) than those observed after topical administration (less than 2 weeks) but are fully reversible. Subcutaneous RTX, however, causes skin ulcerations (that can be severe in some animals) in the head (mostly in the facial skin and around the ears), the cause of which is unclear. It was suggested that RTX may cause paradoxical itch and the ulcers are secondary to the scratching behavior of the animals. This effect of RTX is somewhat different than other vanilloids such as capsaicin and curcumin which reveal anticarcinogenic and antitumor activity (Surh et al. 1995; Park and Surh 1997; Limtrakul et al. 1997; Jang et al. 1989; Huang et al. 1997; Tanaka et al. 2002). Also, high-concentration RTX (leaking from the injection site) may damage the epithelium, either directly (there are reports that keratinocytes may express functional TRPV1 receptors) or indirectly (Li et al. 2007). Finally, intraganglionic RTX injection (into the trigeminal ganglion) was reported to abolish eye wiping in response to capsaicin in an irreversible fashion; this was associated with a loss of TRPV1/CGRP-positive neurons (Karai et al. 2004).

The effect of RTX-induced denervation on TRPV1 expression in surrounding tissues has been recently examined. Surgical removal of both the sciatic and saphenous nerves from rat right hind legs and a parallel setup administration of RTX to the rat leg subcutaneously were performed (Kun et al. 2012). Two weeks after administration, the dorsal and plantar paw skin samples of hind legs, as well as the oral mucosa, were excised. Neither chemical nor surgical denervation influenced the level of TRPV1 receptor mRNA and protein expression in non-neural cells of either skin regions or mucosa (Kun et al. 2012). This indicates that RTX pretreatment is cytotoxic only to TRPV1-positive neurons and does not affect surrounding non-neural tissues.

As discussed above, vanilloid treatment depletes the proinflammatory neuropeptides substance-P (SP) and CGRP from capsaicin-sensitive sensory neurons. It was proposed that capsaicin blocks centripetal intra-axonal transport and thereby starves the cell bodies of nerve growth factor (NGF) which is produced in the periphery and is important for neuropeptide synthesis. Indeed, RTX treatment causes a marked decrease in pre-protachykinin mRNA levels encoding SP (Szallasi et al. 1999). Subsequently, it was discovered that vanilloid-induced changes in neuropeptide levels are, in fact, bidirectional: while SP and CGRP are downregulated after capsaicin or RTX treatment, the expression of an endogenous analgesic peptide galanin is increased. Moreover, increased neuronal nitric oxide synthase (nNOS) and cholecystokinin receptor-B (CCK-B) levels were observed in the RTX-treated animals (Burliński et al. 2011). Collectively, these changes were termed "vanilloid-induced messenger plasticity" (Szallasi and Blumberg 1999). It was postulated that vanilloid treatment alters the phenotype of sensory neurons from proinflammatory to analgesic/anti-inflammatory. Thus, vanilloid-induced messenger plasticity may represent a major mechanism of desensitization.

#### 8 Vanilloids and Mitochondrial Dysfunction

In cultured DRG neurons, there is a clear separation between concentrations at which capsaicin activates TRPV1 ( $ED_{50} < 100 \text{ nM}$ ) and at which it becomes neurotoxic (30–100 µM) (Chard et al. 1995; Wood et al. 1988). Moreover, capsaicin was reported to kill a variety of non-neuronal cells such as in human B-cells, mouse myeloid cell lines, and Jurkat T-cells as well as squamous cell carcinoma cell lines (Wolvetang et al. 1996; Macho et al. 1999; Lee and Surh. 1998; Hail and Lotan. 2002). Capsaicin-induced apoptosis was also described in rat thymocytes (Amantini et al. 2004). It is still hotly debated if these non-neuronal cells express functional TRPV1. For example, in dendritic cells, both the presence and absence of TRPV1 were reported (O'Connell et al. 2005; Tóth et al. 2009).

The effect of capsaicin on cell death seems to be either specific (TRPV1 mediated) or nonspecific (not mediated by TPV1) in nature. Capsaicin can cause apoptosis or necrosis depending on the dose applied. In cultured rat DRG cells, capsaicin causes apoptosis by increasing the intracellular Ca<sup>2+</sup> concentration, enhancing mitochondrial Ca<sup>2+</sup> accumulation, dissipation of the inner transmembrane potential ( $\Delta \psi_m$ ), activation of Ca<sup>2+</sup>-sensitive proteases, and DNA fragmentation (Dedov et al. 2001). Capsaicin evokes similar signalling events in transformed and mitogen-activated T-cells (Macho et al. 1999) and in human and rat glioblastoma cells (Bíró et al. 1998; Lee et al. 2000).

The intercellular Ca<sup>2+</sup> homeostasis is maintained by Ca<sup>2+</sup>-binding proteins present in cytoplasm, endoplasmic reticulum (ER), and mitochondria (Kostyuk and Verkhratsky 1994; Svichar et al 1997; Verkhratsky and Petersen 1998). The elevated intracellular Ca<sup>2+</sup> triggered by capsaicin leads to activation of Ca<sup>2+</sup>-dependent enzymes such as different phospholipases, proteases, and endonucleases that can cause apoptosis in neuronal cells as well as non-neuronal cells (McConkey and Orrenius 1996; Wood et al. 1988). Though in general mitochondria are able to sequester intracellular Ca<sup>2+</sup>, excess Ca<sup>2+</sup>-influx into mitochondria causes membrane permeability transition (MPT) pore in mitochondrial membrane (Wong et al. 2012). This is considered as prototypical inducing factor. This MPT allows water and other small molecules to infiltrate inside the mitochondrial matrix which leads to osmotic swelling of mitochondria and may cause physical rupture of mitochondrial membrane (Green and Reed 1998; Bernardi 1992; Crompton 1999; Kroemer and Reed 2000).

However, several reports suggest that vanilloids may exert effects which are independent of Ca<sup>2+</sup>-influx and TRPV1 receptors. This is due to the fact that capsaicin has significant effects in biological systems that are much lower in the phylogenetic tree and do not contain TRPV1. For example, capsaicin acts as inhibitor for organisms such as Paracoccus denitrificans, Escherichia coli, and Thermus thermophilus HB-8 where it affects ubiquinone reduction by NADH (Yagi 1990). Not only pure capsaicin, other vanilloids such as dihydrocapsaicin and RTX can also act as inhibitor of NADH oxidase (Wolvetang et al. 1996). Indeed, these inhibitors are able to induce apoptosis in human B-cell and mouse myeloid cell line (Wolvetang et al. 1996). In contrast, it was shown that in organisms which do not have the energy transducing site, such as in Saccharomyces cerevisiae mitochondria and Bacillus subtilis membranes, capsaicin does not inhibit NADH-ubiquinone reductase, suggesting that mitochondria can be a potential target of capsaicin action (Yagi 1990). In addition to mitochondria, recently it has been shown that in hippocampal astrocytes capsaicin can act as an inhibitor of tyrosyl tRNA synthetase and by inhibiting this enzyme, it induces cell death (Cochereau et al. 1996, 1997).

The TRPV1-independent functions of capsaicin mostly indicate the deleterious effect of capsaicin on mitochondria. Earlier research suggests that after the systemic application of capsaicin in the A $\beta$ -type sensory neuron of adult rat as well as neonatal rat leads to mitochondrial swelling and results in the formation of atypical hollow mitochondria (Joó et al. 1969; Szolcsányi et al. 1975; Jancsó et al. 1977; Jancsó and Király 1981; Szöke et al. 1998, 2002). However, the real molecular mechanism behind the formation of hollow mitochondria is not clear. In addition to the TRPV1 receptor-mediated effects, it seems that capsaicin can also exert receptor-independent effects on the mitochondria. Due to its structure, capsaicin can act as analog of coenzyme Q, a lipophilic mobile electron carrier present in plasma membrane and involved in maintaining the redox potential of membrane. Indeed, it has been reported that preincubation of human lymphoblastoid cells with coenzyme Q prevents capsaicin-induced apoptosis (Wolvetang et al. 1996; Macho et al. 2000). It suggests that capsaicin competes for coenzyme Q and alters the redox potential of plasma membrane. Apart from that it has been also reported in transformed and activated T-cells; capsaicin inhibits the plasma membrane NADH oxidoreductase (PMOR), an enzyme that transfers electrons from cytoplasmic NADH to external electron acceptors such as oxygen via coenzyme Q (ubiquinone) (Morré et al. 1995, 1996; Wolvetang et al. 1996; Macho et al. 2000). Capsaicin can also inhibit the NADH:coenzyme Q oxidoreductase (complex I) activity of the mitochondrial electron transport system (Shimomura et al. 1989; Yagi 1990) which causes alteration on the mitochondrial membrane structure and its function (Aranda et al. 1995; Tsuchiya 2001). In vitro experiments in transformed cells as well as in activated T-cells suggest that capsaicin treatment enhances the generation of reactive oxygen species (Macho et al. 1998, 1999; Garle et al. 2000) and depolarization of mitochondrial membrane (Dedov et al. 2001) and apoptosis. Capsaicin suppresses the growth of cancer cells by NF-kB inactivation, reactive oxygen species (ROS) generations, cell-cycle arrest, and modulating EGFR/HER-2 pathways (Hail 2003; Kang et al. 2003; Lee et al. 2004; Min et al. 2004; Surh 2002; Thoennissen et al. 2010). Similarly, RTX was also reported to inhibit the NADH oxidase located in plasma membranes, to generate reactive oxygen species, and to induce apoptosis in transformed cells (Wolvetang et al. 1996; Macho et al. 1999, 2000; Garle et al. 2000). The exact molecular mechanism by which capsaicin causes oxidative stress and apoptosis remains rudimentary (Pramanik et al. 2011).

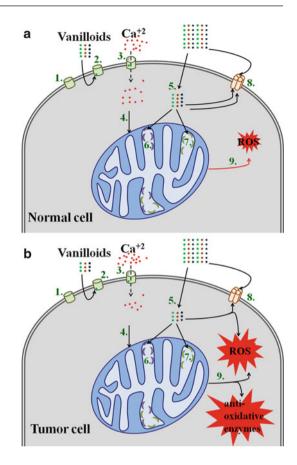
#### 9 Conclusion

Historically, capsaicin and RTX proved to be invaluable tools in defining a fundamental subdivision of the peripheral nervous system (Appendino and Szallasi 1997; Buck and Burks 1986). Capsaicin-sensitive neurons are peptidergic nociceptive (primary sensory) neurons that express the vanilloid (capsaicin) receptor TRPV1. The activation of these neurons not only transmits painful stimuli to the CNS but also initiates the neurogenic inflammatory response. In turn, neurogenic inflammation was postulated to play a pivotal role in the pathogenesis of diverse diseases states, ranging from migraine through asthma to irritable bowel disease.

What makes capsaicin unique among naturally occurring irritant agents is that the initial stimulation of these neurons is followed by a lasting but fully reversible refractory state (traditionally termed desensitization) or irreversible neurotoxicity. Desensitization to capsaicin has a clear therapeutic potential. Indeed, capsaicincontaining creams and patches have clinical application for decades for indications like diabetic polyneuropathy. Irreversible neurotoxicity can also be exploited for therapeutic purposes. Indeed, at present site-specific RTX injections are undergoing clinical trials to achieve permanent analgesia in cancer patients with chronic, intractable pain secondary to metastatic disease.

Despite the tremendous progress that has been made after the identification and molecular cloning of TRPV1 in understanding capsaicin mechanisms, several outstanding questions remain unsolved. Most important, the exact molecular mechanisms underlying reversible desensitization versus irreversible toxicity are yet to be established. At high concentrations, the specificity of vanilloids for TRPV1 is lost, and the distinction between specific (TRPV1-mediated) and nonspecific (non-TRPV1-mediated) vanilloid actions becomes problematic. As an added complication, the existence of functional TRPV1 receptors in cells other than capsaicinsensitive primary sensory neurons remains controversial. For example, TRPV1 knockout mice show altered behavior (e.g., reduced fear response) and capsaicin causes extensive neurodegenerative changes in the rat brain; yet, reporter mice show essentially no TRPV1 expression in the brain (Cavanaugh et al. 2011; Marsch et al. 2007; Tóth et al. 2005; Roberts et al. 2004; Starowicz et al. 2008; Goswami et al. 2010). In this context it is worth to mention that involvement of TRPV1 in the regulation of spine morphology and synaptic transmission has been demonstrated (Goswami et al. 2010). Another puzzling example is the keratinocytes present in skin. Keratinocytes were reported to express TRPV1 (both mRNA and protein) and

Fig. 1 TRPV1-dependent and TRPV1-independent vanilloid actions in normal cell (**a**) and tumor cell (**b**). TRPV1 present in the plasma membrane can be activated by different vanilloids with different affinities and binding kinetics (Steps 1-2). Potent vanilloids such as capsaicin or RTX activate TRPV1 at very low concentration and cause Ca2+ influx (Step 3). Vanilloids at much higher concentrations can cross plasma membrane and can also act on electron transport chain complex 1 (ETC1) and electron transport chain complex 3 (ETC3) (Steps 5-7). Vanilloids can also act on the coenzyme O and PMOR (Steps 8-9). All these factors result in production of ROS (Step 10) relevant for neurotoxicity



respond to capsaicin with  $Ca^{2+}$ -uptake (Southall et al. 2003; Lee et al. 2008; Pecze et al. 2008). Yet, capsaicin evokes no responses in the rat skin after skin denervation (Fig. 1).

The mechanisms responsible for the marked species-related differences in vanilloid actions are only partially understood. It is now clear that birds do not respond to capsaicin because the avian TRPV1 receptor lacks the functional vanilloid-binding motif of mammalian TRPV1 receptors. However, it remains a mystery why the pulmonary chemoreflex is differentially regulated by RTX in rats (desensitization without prior excitation) and cats (repeatable excitation with no desensitization). However, much more and detailed studies are needed to characterize vanilloids for therapeutic applications in near future.

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Part II

## **Dopamine Chemistry and Parkinson Disease**

# 6-Hydroxydopamine as Preclinical Model of Parkinson's Disease

Maria F. Galindo, Sara Saez-Atienzar, Maria E. Solesio, and Joaquín Jordán

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M.F. Galindo (🖂)

Unidad de Neuropsicofarmacología Traslacional, Complejo Hospitalario Universitario de Albacete, Albacete, Spain

Pharmaceutical Technology School of Pharmacy, University of Castilla-La Mancha, Albacete, Spain

e-mail: mgalindoa@sescam.jccm.es

S. Saez-Atienzar

Faculty of Medicine and Dentistry, Catholic University of Valencia San Vicente Mártir, Valencia, Spain

M.E. Solesio

Unidad de Neuropsicofarmacología Traslacional, Complejo Hospitalario Universitario de Albacete, Albacete, Spain

J. Jordán (⊠) Grupo de Neurofarmacología, Facultad de Medicina de Albacete, Universidad de Castilla-La Mancha, Albacete, Spain e-mail: joaquin.jordan@uclm.es

#### Abstract

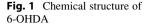
Like many other diseases, Parkinson's disease (PD) requires preclinical models to understand the underlying molecular and cellular mechanisms that cause the disorder. Such models will also help to identify novel potential drug targets and to elucidate the effects of new drug candidates. The hydroxylated analogue of dopamine, 6-hydroxydopamine (6-OHDA), has been exploited as an experimental model to study PD. It also has been used extensively as a test system for novel symptomatic agents and for assessment of neuroprotective and neurorepair strategies. The 6-OHDA lesioned rat appears to be a good predictor for monitoring the efficacy of new dopaminergic drugs that enter phase II/III clinical trials. The aim of this chapter is the molecular and cellular mechanisms that are involved in the preclinical model of Parkinson's disease based on 6-OHDA. With special attention, we address the role of second messengers as the role of reactive oxygen species and the mitochondria as the headquarters of cell death. The role of molecular signaling pathways, for instance, the participation of Bcl-2 family members, will also be addressed.

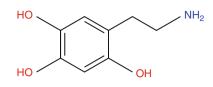
List of Abb	reviations
6-OHDA	6-hydroxydopamine
Bak	Bcl-2-antagonist/killer
Bax	X Bcl-2-associated protein
DA	Dopamine
GSH	Glutathione
MCC	Multiple conductance channel
MOMP	Permeabilization of the mitochondrial outer membrane
PD	Parkinson's disease
PTP	Permeability transition pore
ROS	Reactive oxygen species
$\Delta \Psi m$	Mitochondrial membrane potential

#### 1 Introduction

6-Hydroxydopamine (6-OHDA), also known as oxidopamine, or 2,4,5trihydroxyphenethylamine ( $C_8H_{11}NO_3$ ) is a toxic oxidative metabolite of dopamine and is detected in the brains and urine of Parkinson's disease (PD) patients (Fig. 1). It has been applied broadly to generate experimental models of Parkinson's disease. 6-OHDA shares some structural similarities with dopamine and norepinephrine and exhibits high affinity for several catecholaminergic plasma membrane transporters, such as the dopamine and norepinephrine transporters. Consequently, 6-OHDA can enter both dopaminergic and noradrenergic neurons.

In this chapter we will focus on molecular and cellular mechanisms that are involved in neurodegenerative processes described in PD. We will discuss the preclinical model of Parkinson's disease (based on 6-OHDA), with special attention





to the role of reactive oxygen species as second messengers and mitochondria as the headquarters of cell death. The role of molecular signaling pathways, for instance, the participation of Bcl-2 family members, will also be addressed.

#### 2 In Vivo Experimental Animal Model

6-OHDA poorly crosses the blood-brain barrier, and, therefore, it does not accumulate within the brain parenchyma to meaningful neurotoxic concentrations following systemic injections. Thus, when 6-OHDA is administrated by systemic injection, it will not produce nigrostriatal lesions. Because of this, systemic injection is the preferred way of 6-OHDA administration to damage the peripheral nervous system and cause chemical sympathectomy. 6-OHDA has been found to deplete peripheral organs of noradrenaline (Porter et al. 1963; Laverty and Sharman 1965; Thoenen and Tranzer 1968). Former studies reported that 6-OHDA caused selective, acute degeneration of symptomatic nerve terminals in the peripheral nervous system (Malmfors and Sachs 1968; Thoenen and Tranzer 1968).

Already back in 1968, Thoenen et al. postulated that 6-OHDA can serve as a valuable tool for functional studies of the central nervous system (Thoenen and Tranzer 1968). The injection of 6-OHDA into the substantia nigra and the subsequent depletion of the nigro-neostriatal dopamine (DA) system were associated with marked motor asymmetry in the animals. This is in good agreement with data of Pottier et al. (1966). They found hypokinesia of the limbs ipsilateral to the depletion of striatal amines after the induction of lesions in monkeys. The same type of electrothermal lesion in rats also depleted natal DA (Hokfelt and Ungerstedt 1969). Moreover, it resulted in ipsilateral turning behavior, which is very similar to that found after nigral injection of 6-OHDA. Motor asymmetries were also reported upon removal of the corpus striatum and treatment with various drugs that interfere with catecholamine neurotransmission. Similar data to these were obtained after unilateral injection of DA into the nucleus caudatus putamen, which resulted in contralateral turning behavior (Ungerstedt 1968). From the results listed above, it seems probable that intracerebrally injected 6-OHDA is able to induce degeneration of catecholamine cell bodies, axons, and terminals in the vicinity of the site of injection. It thus offers the possibility of tracing these neuron systems in the brain, for instance, by studying the accumulation and disappearance of the transmitter after local injections of 6-OHDA into various cell groups, axon bundles, or terminal areas. Degeneration of the nigro-neostriatal DA neuron system, caused by 6-OHDA, produces marked motor disturbances, which are known to be associated with this system.

Several local sites of injection have been used to damage the central dopaminergic pathways, including intraventricular, intracisternal, and intracerebral tissues. For reviews on the animal model, see Bove et al. (2005), Cenci et al. (2002), and Duty and Jenner (2011).

#### 3 In Vitro Experimental Cell Culture Models

There is accumulating evidence from in vitro and in vivo studies, implicating cell death in the etiology of the 6-OHDA model of PD. For instance, in neuronal (cerebellar granule cells, primary mesencephalic dopaminergic) and nonneuronal (mesencephalic-derived dopaminergic MN9D, NB41, SH-SY5Y, thymocytes, bovine chromaffin cells,) cell cultures, 6-OHDA induces characteristic biochemical, histochemical, and morphological changes that are characteristic for apoptosis (Blum et al. 2001). Furthermore, intracerebral injection of 6-OHDA causes apoptosis of dopaminergic neurons in the substantia nigra (Bove et al. 2005; Duty and Jenner 2011).

#### 4 Cell Death Pathways Activated by 6-OHDA

#### 4.1 Reactive Oxygen Species

Reactive oxygen species (ROS) are important for execution of physiological functions. However, excessive production of ROS is detrimental to the cell. Following an increase in ROS production, the cell's redox equilibrium is shifted to a more oxidized state, affecting both the structure and the function of different molecules. This may lead to specific toxic processes, which compromise the redox status of the cell and can cause cell death. Due to high levels of polyunsaturated fatty acids in their membranes and the relatively low activity of endogenous antioxidant enzymes, cells in the brain are particularly susceptible to oxidative damage. Increased oxidative stress together with a decline in endogenous antioxidants is an important determinant to develop PD. In our model, 6-OHDA initiates neurodegeneration through a combination of oxidative stress and mitochondrial respiratory dysfunction (Fig. 2).

Under physiological conditions, 6-OHDA is rapidly and nonenzymatically oxidized by molecular oxygen to form 1,4-para-quinone and its degradation products (Gee and Davison 1989), along with production of ROS such as hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), superoxide radical ( $\cdot$ O<sub>2</sub><sup>-</sup>), and hydroxyl radical ( $\cdot$ OH<sup>-</sup>). Quinones react with nucleotic groups of macromolecules, leading to inactive or destroyed quinoproteins. These do not seem to contribute significantly to the observed cytotoxic effects of 6-OHDA. H<sub>2</sub>O<sub>2</sub> can enter the cells and reacts with trace metals to form highly reactive  $\cdot$ OH<sup>-</sup> (Koppenol 2001), which can oxidatively damage proteins, lipids, and DNA (Beckman and Ames 1997). We have shown that 6-OHDA concentrations that were nontoxic to cell cultures did not significantly

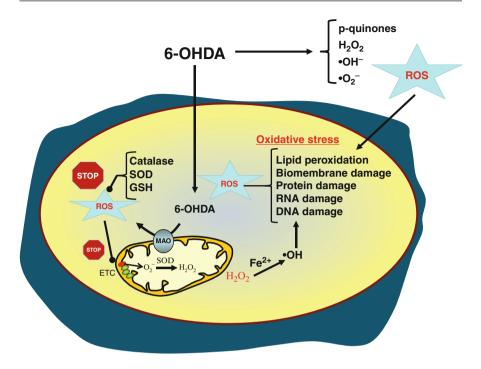


Fig. 2 6-OHDA initiates neurodegeneration through oxidative stress

increase  $H_2O_2$  production (Galindo et al. 2003). Moreover,  $H_2O_2$  addition to cultures produced a pattern of cell death similar to 6-OHDA.

In addition to the nonenzymatic self-auto-oxidation process, microinjection of 6-OHDA into the striatum may lead to the generation of  $H_2O_2$  via a mitochondrial enzymatic oxidation process. Inhibition of complex I of the electron transport chain also stimulated mitochondrial production of superoxide radicals. These superoxide radicals were then catalyzed to  $H_2O_2$  by superoxide dismutase, and, subsequently,  $\bullet OH^-$  may arise from the breakdown of  $H_2O_2$ . This may be associated with the mitochondrial dysfunction seen in our experiments, because  $\bullet OH^-$  rapidly attacks other biological molecules. The radicals produced in molecules such as lipids and proteins may also interact with mitochondrial enzymes to cause degradation.

#### 4.2 Mitochondrial Dysfunction

Mitochondrial dysfunction is associated with many brain diseases including Parkinson's disease, Alzheimer's disease, and brain ischemia (Galindo et al. 2010; Jordan et al. 2011). However, the enzymatic basis for mitochondrial dysfunction remains largely unknown. Mitochondrial dysfunction in PD has been demonstrated as a reduction in complex I activity in the brains of patients (Mizuno et al. 1989; Schapira et al. 1989) and laboratory animals (Dabbeni-Sala et al. 2001).

Increased lipid peroxidation in the substantia nigra is typical of patients with PD (Dexter et al. 1989). This suggests that ROS are produced in excess, resulting in chronic local oxidative stress, which may cause progressive degeneration of the nigral dopaminergic neurons. Normally, ROS are probably produced in large quantities in the substantia nigra of humans, since there is an abundance of detoxifying enzymes, such as superoxide dismutase in dopaminergic neurons (Ceballos et al. 1990) and glutathione peroxidase in glial cells. In parkinsonian patients, the activity of these enzymes is not by definition abnormal (Marttila et al. 1988). However, decreased concentrations of glutathione (Perry et al. 1982) and increased activity of mitochondrial superoxide dismutase (Saggu et al. 1989) indicate that an excess of free radicals may be produced. Importantly, 6-OHDA readily reduces striatal levels of antioxidant enzymes (total glutathione (GSH) or superoxide dismutase) (Kunikowska and Jenner 2001: Perumal et al. 1992) to interact directly with complexes I and V of the mitochondrial respiratory chain. This leads to subsequent respiratory inhibition and further oxidative stress. Many of these effects are thought to mirror events that occur during PD in the brain, and this strongly supports the validity of the 6-OHDA model.

The oxidant  $H_2O_2$  elicits the suppression of thiol-dependent electron transport, which appears to be related to the inhibition of complex I (nicotinamide adenine dinucleotide coenzyme Q reductase) in the respiratory chain (Mizuno et al. 1989). A 1-chloro-2,4-dinitrobenzene reductase in the mitochondrial matrix has been shown to be a thioredoxin reductase that reduces glutathione disulfide to glutathione (Lenartowicz and Wudarczyk 1995). 6-OHDA induced thiol oxidation in brain mitochondria, which was diminished by antioxidant enzymes. DA toxicity on neuronal cells is inhibited by thiol compounds and is potentiated by mitochondrial function inhibitors. Consequently, we and others have observed protection against 6-OHDA-induced toxicity in cell cultures by treatment with different antioxidant compounds (Galindo et al. 2003).

#### 4.3 Mitochondria as Central Regulators of 6-OHDA Neurotoxicity

Mitochondria play critical roles in regulating cellular viability and show characteristic vulnerability to injury. Disruption of mitochondrial  $Ca^{2+}$  homeostasis may be involved in neuronal cell death. Mitochondria exposed to oxidative stress show enhanced  $Ca^{2+}$  uptake and increased formation of ROS. 6-OHDA slightly enhanced the  $Ca^{2+}$  uptake by mitochondria and apparently stimulated ruthenium red-induced  $Ca^{2+}$  release. The inhibitory effect of antioxidant enzymes indicates that ROS may be involved in the mitochondrial  $Ca^{2+}$  elevation induced by catecholamines.

Mitochondria can be considered as headquarters where the cell controls signaling pathways that under some circumstances can lead to cell death (Jordan et al. 2003a, 2011). Mitochondrial membrane permeabilization is a critical event during apoptosis and represents the point of no return of this lethal process (Galluzzi et al. 2009). Permeabilization of the mitochondrial outer membrane (MOMP) is a crucial

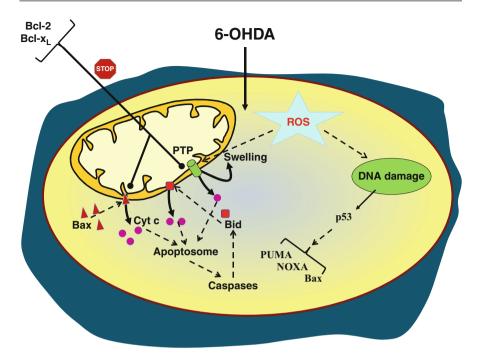


Fig. 3 Hypothetical mechanisms of 6-OHDA-induced MOMP

step both in apoptosis and necrosis. This phenomenon allows the release of mitochondrial death factors, which facilitates or triggers different signaling cascades. Ultimately this causes the execution of the cell involving release of cytochrome c. Treatment with 6-OHDA induced the release of cytochrome c from brain mitochondria (Gomez-Lazaro et al. 2008b).

MOMP is regulated either by the formation of the permeability transition pore (PTP) or by the insertion of Bcl-2 family members into the MOM (Fig. 3). There is some controversy about the role of the PTP in the 6-OHDA model. The PTP is a nonspecific large proteinaceous pore, spanning both mitochondrial membranes. It allows the passage of ions and substrates, in cost of loss of mitochondrial membrane potential. This leads to ATP depletion and energetic collapse and thus contributes to cell death (Qian et al. 1999). Onset of a PTP is such a severe perturbation of mitochondrial function that it basically assures cell death.

The PTP has been proposed to be another target of DA oxidation products in mitochondria. The oxidation of DA to dopamine quinone causes a significant increase in the swelling of brain and liver mitochondria. The effect of dopamine and 6-OHDA on mitochondrial membrane permeability has not been clearly elucidated. We have demonstrated that 6-OHDA did not induce PTP formation (Gomez-Lazaro et al. 2008b). First, 6-OHDA did not decrease the levels of mitochondrial calcein fluorescence, a hallmark of PTP opening. Second, 6-OHDA-induced

mitochondrial cytochrome c release was not inhibited in cell cultures that were pretreated with the PTP inhibitor cyclosporine A. Third, a mitochondria suspension isolated from brain did not readily undergo swelling upon exposure to 6-OHDA.

6-OHDA is also able to activate the multiple conductance channel (MCC), which is located in the inner membrane of mitochondria (Jordan et al. 2004). Normally, under physiological conditions, the MCC is closed, but it can be activated by different stimuli, such as  $Ca^{2+}$  at millimolar concentrations. Activation of the MCC leads to a multitude of changes in mitochondria, including mitochondrial membrane potential ( $\Delta\Psi$ m) dissipation.

6-OHDA induces profound mitochondrial fission in SH-SY5Y cell cultures, an event that precedes collapse of the  $\Delta\Psi$ m and cytochrome *c* release (Gomez-Lazaro et al. 2008a). Mitochondria form a highly dynamic semi-tubular network, the morphology of which is regulated by frequent fission and fusion events as well as by movements along the cytoskeleton (Hoppins et al. 2007). Although the physiological significance of mitochondrial dynamics is not fully understood, growing evidence indicates that maintaining correct mitochondrial morphology through fission and fusion is critical for cell function (Chand et al. 2006; Chen and Chan 2006; Hoppins et al. 2007; Kiefel et al. 2006; Rouiller 1960). A reduction of the mitochondrial volume has been observed in pathological tissue (Arbustini et al. 1998; Nishino et al. 1998).

#### 4.4 Bcl-2 Family Participation

Alternatively, MOMP may be initiated by insertion of the proto-oncogene Bcl-2 family into the MOM. The Bcl-2 family is composed of about 25 key regulators of apoptotic processes. These proteins are structural and functional homologs of the nematode protein CED-9 and are localized in the mitochondrial membrane. They contain up to four regions with a high homology to Bcl-2 (regions BH 1 to 4). Members containing only the BH3 region are proapoptotic proteins, and among them are Bax (X Bcl-2-associated protein), Bak (Bcl-2-antagonist/killer), BIM, and BID. How these proteins modulate changes in mitochondrial permeability is still an enigma. Inactive Bax resides in the cytosol or is anchored to the laxly face of the membranes of various organelles (Wolter et al. 1997). After a cell death signal, the Bax protein acquires a homo-oligomeric shape and is incorporated into the outer mitochondrial membrane. Postmortem studies indicated that the presence of Bax and its translocation to the outer mitochondrial membrane may contribute to the death of dopaminergic neurons in PD (Hartmann et al. 2001).

The protein BID is another Bcl-2 family member that migrates to mitochondria during processes of cell death. This protein is activated by caspase-8 and gives rise to a spoofed form t-bid, which is able to migrate to mitochondria, thereby altering their permeability. This migration is considered as the link between apoptotic caspase-8, triggered by signals of cell death, and downstream signaling cascades. On the other hand, some members of the Bcl-2 family have been proposed to modulate PTP formation. One example is Bcl-2, which inhibits the formation

of the PTP. Bcl-2 stabilizes the mitochondrial membrane, increases the buffering capacity of Ca<sup>2+</sup>, and protects the  $\Delta\Psi$ m to various stimuli. Yamada et al. have demonstrated that injection of a Bcl-2-producing vector into the substantia nigra 1 week prior to 6-OHDA injection increased neuronal survival in this tissue after the lesion (Yamada et al. 1999).

A potent induction of the BH3-only protein PUMA has been observed in response to 6-OHDA at both mRNA and protein levels (Gomez-Lazaro et al. 2008b). PUMA is thought to mediate the majority of the proapoptotic effects of p53 activity, which correlates with neuronal death in neurodegenerative diseases (Gomez-Lazaro et al. 2004; Gomez-Lazaro et al. 2005; Jordan et al. 2003b; Steffan et al. 2000; Tieu et al. 2001) including PD (Duan et al. 2002; Mandir et al. 2002).

Bcl-xL also belongs to the Bcl-2 family of proteins. Bcl-xL is able to protect cells from apoptosis induced by various stimuli, including serum deprivation, heat shock, malonate, and chemotherapeutic drugs (Fernandez-Gomez et al. 2005; Tsujimoto 1989). This suggests that the ability of Bcl-xL to prevent apoptosis is achieved by blocking a common cell death pathway. Our data indicate that Bcl-xL might prevent 6-OHDA effects by inhibiting MCC activation by the toxin (Jordan et al. 2004). TAT-Bcl-xL further reduces the number of cells that contain fragmented DNA or are caspase-3-reactive and increases the number of viable neurons in the striatum (Cao et al. 2002).

#### 4.5 Caspases

Finally, the caspase family of proteases comprises fundamental components of many apoptotics signaling pathways. The percentage of active caspase-3-positive neurons among dopaminergic neurons was significantly higher in PD patients than in controls (Hartmann et al. 2000). Electron microscopy analysis of the human brain and in vitro data suggest that caspase-3 activation precedes and is not a consequence of apoptotic cell death in PD. Activation of caspase-3 was also demonstrated in 6-OHDA-induced apoptosis of SH-SY5Y cells (Gomez-Lazaro et al. 2008b). Pretreatment with caspase inhibitors prevented 6-OHDA-induced apoptosis (Hartmann et al. 2000; Takai et al. 1998). In human PD brain studies, a significantly higher percentage of dopaminergic substantia nigra neurons displayed activation of caspase-1, caspase-3, caspase-8, and caspase-9 (Hartmann et al. 2000; Mogi et al. 2000).

#### 5 Flaws of the Model

However, the predictive validity of this model remains uncertain; strategies that reduced nigral dopaminergic cell loss in experimental animals so far have not been translated to the clinic. This can be due to the fact that the 6-OHDA model does not capture all features of PD. For example, it cannot exclude that PD pathological changes may occur in many brain areas outside of the basal ganglia locus coeruleus and raphe nuclei, and this is not incorporated in the 6-OHDA model. Moreover, in the 6-OHDA model cell death occurs far more rapidly than in PD. Nevertheless, as long as these limitations are understood and accounted for when interpreting effects, the 6-OHDA model will remain at the forefront of preclinical drug discovery for PD.

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### Advances in Stem Cell Research for Parkinson Disease

#### Irmgard Paris, Ulises Ahumada-Castro, and Juan Segura-Aguilar

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#### Abstract

There is significant interest in using stem cells as a cell source for treatment against Parkinson's and other diseases due to the limited self-repair ability of the central nervous system and the lack of a pharmacological treatment to replace lost

I. Paris (🖂)

Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

Department Basic Sciences, Santo Tomas University, Viña del Mar, Chile e-mail: irmgardparis@gmail.com

U. Ahumada-Castro • J. Segura-Aguilar Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

neurons. Parkinson's disease is one of the main neurodegenerative disorders, and it is characterized by the degeneration of the nigrostriatal pathway caused by loss of ventral midbrain dopaminergic neurons in the substantia nigra pars compacta. Currently, there are no treatments to halt disease progress; the treatments merely ameliorate motor symptoms for a time, but the treatments are often followed by severe side effects. Cell-replacement therapy has been proposed as a treatment to replace damaged neurons and protect the remaining cells. Preliminary studies have utilized adult tissues, embryonic/fetal tissues, and embryonic stem cells as a source of dopaminergic cells. However, the ethical issues related to this type of procedure, the high quantity of required cells, the risk of tumor formation, and the immunological rejection associated with it have prevented this technique from being an effective treatment against Parkinson's disease. Researchers have been challenged to focus on new cell-replacement alternatives. In 2006, researchers found that only four transcription factors were necessary to reprogram an adult somatic cell to an induced pluripotent stem cell. This breakthrough revolutionized the scientific community, as this technique would allow scientists to obtain specific cells from the patient, avoiding rejection problems from non-autologous grafts and avoiding ethical problems. In addition, a large number of cells could be obtained. Although induced pluripotent stem (iPS) cells have great potential as a treatment against this disease, there are some obstacles for its clinical use such as the lack of standardized protocols that result in a dopaminergic cell phenotype that grafts without undifferentiated cells and without a risk of tumor formation. Currently, the iPS cell system is an ideal model of study for many other diseases besides Parkinson's, and the system is also good for drug screening. Recently, the direct reprogramming of somatic cells to induce neuronal (iN) cells without generating a pluripotent stage could be the solution to avoiding tumor formation risk. However, as a result of the low quantity of cells obtained, more investigations are necessary to convert this technique into an efficient cell-replacement therapy. If we overcome these obstacles, the use of iPS and iN cells could be, without a doubt, a promising therapy in the near future.

#### Keywords

Dopaminergic neurons • Induced neuronal cells • Induced pluripotent stem cells • Parkinson's disease • Pluripotency • Reprogramming • Stem cell • Therapy

List of Abbreviations				
ASCs	Adult stem cells			
ATRA	All-trans-retinoic acid			
BDNF	Brain-derived neurotrophic factor			
bFGF	Basic fibroblast growth factor			
CB-SCs	Cord blood-derived multipotent stem cells			
CNS	Central nervous system			
DA	Dopamine			
EGF	Epidermal growth factor			

ESCs	Embryonic stem cells
FGF2	Fibroblast growth factor 2
FGF20	Fibroblast growth factor 20
FGF8	Fibroblast growth factor 8
GDNF	•
	Glial cell line-derived neurotrophic factor
GSK-3	Glycogen synthase kinase 3
hESCs	Human embryonic stem cells
HGF	Hepatocyte growth factor
HSCs	Hematopoietic stem cells
HUVMSCs	Human umbilical vein mesenchymal stem cells
ICM	Inner cell mass
iN	Induced neuronal
iPSCs	Induced pluripotent stem cells
MSCs	Mesenchymal stem cells
NP	Neural progenitor
NPSCs	Neural progenitor stem cells
NSCs	Neural stem cells
NSPCs	Neural stem cell and neural progenitor cell
PD	Parkinson's disease
PGCs	Primordial germ cells
SCC	Stem cell coactivator complex
SDF1a	Stromal cell-derived factor-1a
SDIA	Stromal cell-derived inducing activity
sFRP1	Secreted frizzled-related protein 1
SHH	Sonic hedgehog
SNpc	Substantia nigra pars compacta
STEMCCA	Single excisable polycistronic lentiviral stem cell cassette
SVZ	Subventricular zone
VEGFD	Vascular endothelial growth factor D
	vascular endothellar growth factor D

#### 1 Stem Cells

The human body consists of different cell types organized into tissues and organs. In the adult state, tissues are maintained because of the proliferative ability of stem cells (SCs). However, not all tissues have the same potential for renewal.

Different adult tissues differ in terms of the proportion of proliferative cells. For example, proliferative cells were discovered in the pancreas that have been described as being different from stem cells (Dor et al. 2004; Zhao et al. 2008). In the pancreas, the cells were able to renew constantly. Normally, the proliferative capacity of tissue is maintained by stem cells that have not only extensive renewal capacity, but also the ability to generate daughter cells capable of differentiating. These cells are called adult stem cells (ASCs) (Watt and Driskell 2010).

#### 1.1 Characteristics

Stem cells are unspecialized cells in an undifferentiated state that have an extraordinary capacity to proliferate indefinitely without differentiating. However, stem cells are capable of differentiating into one or more specialized cell types. A stem cell divides to generate one daughter stem cell and one progenitor cell during self-renewal; later, the progenitor cell achieves a fully differentiated state.

The stem cells can also be categorized by their ability to divide and produce differentiated cells. These cells can be classified as totipotent, pluripotent, and multipotent cells. Totipotent cells are those that give rise to all cells of an organism including germ cells, as in a zygote. Pluripotent cells generate all cell types in an embryo except the cells of the embryonic membrane such as the placenta. The cells of the inner cell mass (ICM) are pluripotent cells are adult stem cells that, despite the fact that they can give rise to different cell types, only generate a given cell lineage. Adult stem cells can differentiate into more than one mature cell. Examples of these adult stem cells are hematopoietic stem cells, mesenchymal stem cells, and neural stem cells (NSCs).

The stem cells present with different types of differentiation: (i) direct differentiation, where a specific cell type develops in a unidirectional pathway; (ii) transdifferentiation, where a direct conversion occurs from one cell type to another, different, cell type; and (iii) dedifferentiation, where a unipotent stem cell generates a multipotent one. Moreover, a stem cell can also fuse with a somatic cell resulting in another lineage (Nadig 2010; Lunn et al. 2011).

#### 1.2 Classification

Stem cells can also be classified according to their origin: embryonic stem cells (ESCs) and adult stem cells (postnatal stem cells/somatic stem cells). ESCs can be obtained from blastocyst, epiblast, and primordial germ cells. These cells have been described as pluripotent because they are able to generate all cell types of the three primary germ layers (ectoderm, mesoderm, and endoderm). Initially, embryonic stem cells were isolated from preimplantation mouse embryos in the blastocyst state (Evans and Kaufman 1981; Martin 1981). In the year 1998, it was reported that stem cells could also be derived from human blastocysts (Thomson et al. 1998). Pluripotent ESCs can be obtained from the post-implantation epiblasts of the mouse embryo (Brons et al. 2007; Tesar et al. 2007). In this case, ESCs were isolated from 12-day-old embryos called epiblasts. Likewise, pluripotent ESCs can be obtained from primordial germ cells, which are progenitors of adult gametes (Kerr et al. 2006). The use of embryonic tissue in research and clinical fields generates cultural disagreement, as cells are generally obtained from "stored" in vitro fertilization embryos. Ethical and cultural matters make the clinical practice with stem cellbased therapies difficult (Watt and Driskell 2010).

In contrast, ASCs are found in most adult tissues. Unlike ESCs, ASCs are not pluripotent cells but rather multipotent cells. ASCs have the ability to differentiate into more than one cell type but not all cell types, as a pluripotent cell does. Examples of such cells are hematopoietic stem cells (HSCs) and mesenchymal stem cells (MSCs). Hematopoietic stem cells can be obtained from cord or peripheral blood. These stem cells have been widely used in the treatment of several diseases such as leukemia, immunodeficiency, metabolic disorders, and tumors (Reimann et al. 2009; Perl et al. 2010). Mesenchymal stem cells originate in the stromal component of the bone marrow, although it is also possible to derive them from peripheral and umbilical cord blood (Perl et al. 2010). Mesenchymal stem cells are found in the bone marrow and hepatic, dermal, orofacial, and adult dental pulp tissues, among others (Gronthos et al. 2000; Nadig 2009; Nesti et al. 2011). Consequently, the use of MSCs as a treatment has been investigated for several diseases (Mundra et al. 2013; Morigi and Benigni 2012; Vawda and Fehlings 2012). Likewise, the role of MSCs in the contribution to tumor growth and progression has also been studied (Yang et al. 2013).

Currently, several human stem cell lines have been characterized that exhibit similar expression patterns for several pluripotency and lineage markers. However, these lines also present differences mainly due to the techniques used for their derivation and maintenance. Consequently, epigenetic factors and intrinsic genetic variations are important factors inherent to human samples, which could contribute to these variations and thus affect gene expression patterns (Adewumi et al. 2007).

#### 1.3 Generation of Pluripotency

In the past, it was believed that only ESCs are pluripotent. However, further investigations have shown that not only are ESCs pluripotent, but adult cells can, in some circumstances, also produce progeny that differentiate across the three primary germ layers (ectoderm, mesoderm, and endoderm), which makes them pluripotent. A mechanism of reprogramming an adult cell to a pluripotent state is the transference of an adult nucleus into the cytoplasm of an oocyte (Gurdon et al. 1958; Gurdon and Melton 2008). An example of this technique is the creation of Dolly the sheep (Wilmut et al. 1997). Likewise, another mechanism to generate pluripotent cells is the fusion of adult cells with a pluripotent cell (Miller and Ruddle 1976). As described above, a pluripotent state can be generated through somatic nuclear transfer. Adult cells can also be reprogrammed to a pluripotent state by the transfection of only four transcription factors: Oct4, Sox2, Klf4, and Myc (Takahashi and Yamanaka 2006).

#### 1.4 Markers of Pluripotency

During early embryonic development, cells change their gene expression profile. Some pluripotent state-related genes are expressed, whereas those related to differentiation should be transcriptionally silenced. Many studies have identified the diverse pluripotency factors that are expressed in both embryos and stem cells.

Several methods have allowed for the identification of pluripotency factors. Studies performed in embryos, cell lines, and ESCs have relied on candidate gene analysis. Phenotypes resulting from specific knockdown, RNAi screens (Chambers et al. 2003; Rodda et al. 2005; Ivanova et al. 2006), gene expression profiles (Voutila et al. 2012; Zheng and Hu 2012), epigenetic profiles (Tomkins et al. 2012; Seisenberger et al. 2012; Al-Khtib et al. 2012), and protein expression profiles (Loh et al. 2006; Walker et al. 2007) have extended our knowledge of the genes and molecules associated with pluripotency.

These studies have allowed for the description of the role of the transcription factors Oct4, Sox2, Klf4, c-Myc, and Nanog for the establishment and/or maintenance of the pluripotent state.

However, a substantial amount remains to be learned regarding the changes in gene expression that occur during embryo development. Such changes are responsible for the pluripotent stem cell state and later to cell differentiation. In addition, it is crucial to understand the role of exogenous and endogenous factors for stem cell maintenance and how, through signaling pathways, gene expression levels are impacted.

These studies have helped determine a list of pluripotency-related genes that might be possible candidates for cellular reprogramming. However, the studies have not clarified the differential expression of genes leading to the pluripotent stem cell state nor how these genes determine the maintenance of the pluripotent state. It is important to emphasize that signaling pathways controlling pluripotency are extensively complex.

#### 2 Stem Cells and Parkinson's Disease

#### 2.1 Parkinson's Disease

Parkinson's disease (PD) is a neurodegenerative disorder that affects approximately 1 % of the population over 65 years of age worldwide (Twelves et al. 2003). However, the prevalence of this disease increases with the age. Studies performed in the USA have shown that PD incidence varies among Whites, Blacks, and Asians, and PD is substantially more common in Whites (Wright Willis et al. 2010). PD was described for the first time by Dr. James Parkinson in the year 1817. Although the disease etiology remains unknown, several factors are believed to be involved in PD development. This disease involves widespread dysfunction and cell degeneration that is characterized mainly by the gradual degeneration of ventral midbrain dopaminergic neurons in the substantia nigra pars compacta (SNpc), causing nigrostriatal pathway dysfunction. This degeneration leads to concomitant depletion dopamine (DA) in the striatum corpus, mostly in the caudate and putamen areas (Olanow 2004). This nigrostriatal pathway is related to control-ling movement and gait. It should be noted that many patients suffering from

Parkinson's disease experience not only motor symptoms but also non-motor ones (Storch et al. 2013).

Parkinson's disease is mainly diagnosed by the motor clinical manifestations presented by patients. Unfortunately, the first motor symptoms are normally observed when more than 60 % of the dopaminergic neurons have died (Bernheimer et al. 1973). Currently, strong evidence shows the presence of non-motor symptoms such as olfactory dysfunction, sleep abnormalities, cardiac sympathetic denervation, depression, and pain long before the presence of motor symptoms (O'Sullivan et al. 2008; Obeso et al. 2010; Storch et al. 2013). These findings are particularly encouraging because they could aid in disease diagnosis at an early stage and thus result in therapies to halt its progress.

Currently, there are no effective treatments to cure this disease, and pharmacological therapies with dopaminergic drugs or functional neurosurgery merely improves the motor symptoms. The pharmacological treatments include dopamine precursors or dopamine agonists. It has been noted that treatment with L-dopa, the precursor of dopamine, may alleviate symptoms for only a few years. Moreover, long-term treatment with L-dopa is not satisfactory because the remaining dopaminergic neurons continue degenerating (Marsden 1994). In addition, chronic drug use is related to a variety of adverse effects such as L-dopa-induced dyskinesias (Politis et al. 2003; Fisone and Bezard 2011). Likewise, deep brain stimulation directed to the subthalamic nucleus or globus pallidus also may relieve symptoms in some patients (Odekerken et al. 2013). Current treatment options are oriented to reverse motors features of this disease such as tremor, rigidity, and bradykinesia/ akinesia; however, these treatments do not halt PD progress. As the main cause of symptoms is ongoing degeneration, new treatments should, in the future, focus on halting dopaminergic neuron degeneration using neuroprotective agents or cell therapies to replace the degenerated neurons. Therefore, replacement therapy with stem cells or fetal tissue might be the best approach (Pawitan 2011), although many issues remain to be resolved before this technique can be used therapeutically. Transplanted cell survival is a major obstacle for the development of cell therapies for PD.

#### 2.2 Stem Cell Therapy for Parkinson's Disease

Currently, PD patients are treated with various drugs including L-dopa, dopaminergic agonists, and inhibitors of dopamine-degrading enzymes. Likewise, deep brain stimulation has been utilized in numerous patients. While these treatments are effective early on to ameliorate the motor symptoms, the treatments do not halt the disease course. In recent decades, research efforts have focused on replacing the dopaminergic neurons that are lost in PD patients. To this end, cellular therapies that replace degenerated cells are the main interest focus today. These therapies involve the use of tissue transplants such as fetal, embryonic, or adult tissues as a source of dopaminergic cells or neural progenitor cells (Lunn et al. 2011). Over time, investigations included not only transplants from the same species but also used autotransplantation and xenotransplantation between different species such as rats, mice, monkeys, pigs, and humans (Huffaker et al. 1989; Bankiewicz et al. 1990; Freed et al. 1990; Galpern et al. 1996; Hara et al. 1997; Molenaar et al. 1997; Barinaga 2000; Yurek and Fletcher-Turner 2004; Puschban et al. 2005; Arias-Carrión and Yuan 2009). Furthermore, investigations have performed the direct transplantation of SC-derived dopaminergic cells such as those derived from ESCs, induced pluripotent stem cells (iPSCs), fetal brain NSCs, and bone marrow-derived SC (Kim 2011). As recently described, other cell types, different from stem cells, may also be suitable for cell therapy. These cells include the induced neuronal (iN) cells, which are derived from somatic cells (Caiazzo et al. 2011; Politis and Lindvall 2012; Lévêque et al. 2012). Moreover, innovative therapies have emerged involving transplantation with genetically modified neuronal or nonneuronal cells. In some cases, the cells were genetically manipulated by introducing genes such as tyrosine hydroxylase, Bcl2, Nurr1, or neurotrophic factors (Holm et al. 2001; Torres et al. 2005; Shim et al. 2007; Ramos-Moreno et al. 2012).

One of the first transplantation studies was performed using a brain tissue graft containing dopamine-secreting cells in the rat striatum. The results were encouraging because the motor abnormalities present in these rats were significantly reduced after the fetal brain tissue graft (Perlow et al. 1979). Subsequent studies proceeded with searching for new sources of donor dopaminergic cells; some studies were very promising but others had poor results. The adrenal medulla, fetal or embryonic mesencephalic tissue, and the carotid body have been used as a source of dopaminesecreting cells (Strömberg et al. 1986; Espejo et al. 1998). Transplantation of human adrenal chromaffin cells or sympathetic neurons from the cervical ganglion was not effective (Yong et al. 1989). Adrenal medulla transplants also showed inconsistent results. Despite the fact that co-transplantation of peripheral nerves and the adrenal medulla in patients with Parkinson's disease can promote long-term improvement, other studies performed in patients showed that the adrenal medullary transplants do not survive long-term (López-Lozano et al. 2000; Kompoliti et al. 2007). On the other hand, the transplantation of mesencephalic dopaminergic neurons or the intrastriatal transplantation of carotid body tissues show more promise as future therapies for this disease (Mendez et al. 2005; Mínguez-Castellanos et al. 2007; Pardal and López-Barneo 2012).

The carotid body consists of adult tissue that contains dopamine-secreting cells and expresses high levels of glial cell line-derived neurotrophic factor GDNF (Toledo-Aral et al. 2003; Porzionato et al. 2008). Studies have shown that intrastriatal autotransplantation of carotid tissues induced functional recovery in parkinsonian rats and monkeys and in some patients with advanced PD (Espejo et al. 1998; Luquin et al. 1999; Arjona et al. 2003). More recently, a population of neural progenitor stem cells (NPSCs) was discovered in this organ (Pardal et al. 2007). Hence, not only is the transplantation of dopaminergic cells necessary for cell therapy, but also the presence of neurotrophic factors as a neuroprotection mechanism. These studies suggest the use of neurotrophic factors as a new experimental approach to combat this disease (Minguez-Castellanos and Escamilla-Sevilla 2005). There is concern because the grafted cells not only produce dopamine, but also deliver neurotrophic factors that protect nigrostriatal neurons. Thus, the trophic factors would be capable of promoting restoration of the nigrostriatal pathway. In this regard, patient-specific adult stem cells derived from the carotid body could be an interesting candidate for PD treatment that avoids human embryonic/fetal tissue or embryonic stem cell use. The cells derived from adult tissues could also be expanded and differentiated into DA precursor cells in vitro (Pardal and López-Barneo 2012). More recently, studies of co-grafts of a cell suspension of rat dopaminergic neuroblasts with carotid body cells increased survival of dopaminergic neurons in rats with total unilateral dopaminergic denervation (Rodriguez-Pallares et al. 2012).

In spite of the fact that cell therapies with carotid tissues appeared to be effective in animal models and patients, the amount of tissue necessary for obtaining significant clinical benefit is one of the main factors limiting their use clinically. Although we are able to develop optimized protocols that could improve the safety and efficacy of these procedures, a major constraint is the availability of tissue for these grafts.

In this respect, progress in stem cell research appears to be the future for cell therapy against Parkinson's disease. Due to the self-renewal capacity and multilineage plasticity of stem cells, stem cells can generate numerous cell types needed for cell therapy. As mentioned previously, several types of stem cells have been isolated from many sources such as embryonic tissues, peripheral blood, umbilical cord blood, bone marrow, and the olfactory bulb. Stem cells have also been obtained from somatic cells; these stem cells comprise the ESCs, iPSCs, NSCs, and MSCs.

#### 2.2.1 Embryonic/Fetal Tissue Cell Therapy for Parkinson's Disease

In the year 1988, Brundin and collaborators performed the first transplantation of human fetal mesencephalic neurons. The results were promising, showing that patients can effectively restore dopaminergic neurotransmission (Brundin et al. 1988). Because the utilization of human fetal tissue as source of dopamine cells is not easy, xenografting of the pig ventral mesencephalon may be an appropriate alternative to the use of human fetal tissue as a PD therapy. These studies showed how implants could survive and mediate functional recovery in animal model of PD and also in some patients, although immunological rejection is the main limitation in humans (Huffaker et al. 1989; Bankiewicz et al. 1990; Galpern et al. 1996; Molenaar et al. 1997; Schumacher et al. 2000; Larsson and Widner 2000).

One of the most commonly used therapies has been the intrastriatal transplantation of embryonic or fetal mesencephalic neurons, which have produced the best clinical results (Madrazo et al. 1990; Freed et al. 1990, 1992, 2001; Sirinathsinghji et al. 1990; Lindvall et al. 1994; Wenning et al. 1997; Piccini et al. 1999). Although in some cases the implanted neuron can survive many years, this phenomenon was not observed in all patients (Piccini et al. 1999). Thus, one of the main difficulties observed with these grafts is the low survival of implanted neurons (Piccini et al. 1999; Emgård et al. 1999; Kaminski Schierle et al. 1999). For this reason, further therapies were oriented to the application of neurotrophic factors as a neuroprotective mechanism of neuronal degeneration (Kupsch et al. 1995; Ballagi et al. 1994; Collier et al. 1999). On the other hand, to improve transplant efficiency, new alternatives emerged including double grafts, either sequentially transplanted or co-grafted (Mendez et al. 1996; Sortwell et al. 1998; Hagell et al. 1999; Kaddis et al. 2000). Simultaneous intrastriatal and intranigral neuronal transplants in rat lesions contribute to striatal reinnervation and reestablishment of the nigrostriatal circuitry (Mendez et al. 1996). Moreover, sequential transplantation of human embryonic mesencephalic tissue in PD patients showed restoring of fluorodopa uptake without affecting graft survival (Hagell et al. 1999). Interestingly, co-grafts of DA neurons with Schwann cells, muscle cells with mesencephalic tissue, or mesencephalic tissue with striatal cell suspensions increase the survival and outgrowth of the DA neurons present in mesencephalic tissue. These effects were related to the neurotrophic factors delivered by the Schwann, skeletal muscle, and striatal cells (Sortwell et al. 1998; Collier et al. 1999; Kaddis et al. 2000). An interesting study performed in 250 patients with advanced PD showed how transplanted embryonic neuronal grafts survive and induce significant therapeutic effects (Björklund and Lindvall 1999).

All investigations performed in animals and humans have contributed to establishing the requirements of the transplanted cells to be used as a cell therapy for Parkinson's disease. Therefore, some important points to consider regarding establishing transplanted cells as possible cell therapy are the following: the presence of functional and morphological characteristics similar to dopaminergic neurons; the existence of the capacity to reestablish nigrostriatal circuitry; the presence of greater cell survival in tissue; the absence of adverse effects such as transplant-associated dyskinesias, transplant rejection, or tumor formation; and the reversion of motor symptoms in PD patients and animal models. On the other hand, difficulties in obtaining fetal tissue have added to the adverse effects caused by transplantation (the graft-induced dyskinesias in some patients), and the low survival of grafted cells makes this technique unsuitable for practice in numerous patients and cannot be recommended as a PD therapy (Freed et al. 2001; Olanow et al. 2003). In this regard, other neuronal cell sources such as embryonic stem cells could be an alternative cell therapy for PD.

## 2.2.2 Embryonic Stem Cell Therapy for Parkinson's Disease

Embryonic stem cells can be derived from different embryonic developmental states such as blastocyst, epiblast, and also primordial germ cells (Evans and Kaufman 1981; Martin 1981; Thomson et al. 1998; Kerr et al. 2006; Brons et al. 2007). These ESCs possess a self-renewal ability and retain pluripotency over prolonged periods of expansion in culture without differentiation (Dushnik-Levinson and Benvenisty 1995; Amit et al. 2000). Unlike other cell types, ESCs have a higher plasticity and the capacity to differentiate into any cell type including dopamine neurons (Park et al. 2004; Zeng et al. 2004; Yan et al. 2005; Datta et al. 2013). One of the main limitations of using ESCs for transplantation in PD patients is the pluripotent state of these cells because under these conditions, the cells could form tumors once grafted, hindering their clinical application. To avoid adverse

effects, an early study showed that an enriched population of midbrain NSCs could be derived from ESCs. Thus, the dopamine neurons obtained from neural SCs are suitable for transplantation (Kim et al. 2002). A recent report showed that dopamine neurons derived from human ESCs efficiently survive after transplantation in animal PD models without forming tumors (Kriks et al. 2011).

The decreased survival of grafted cells after transplantation is one of the main obstacles for this surgery. One procedure involves the injection of adenoviral vectors into the dopamine-depleted striatum of hemiparkinsonian rats to deliver the differentiation factor sonic hedgehog (SHH) or the glial cell line-derived neurotrophic factor (GDNF) to dopamine-rich grafts. Later, embryonic ventral mesencephalic cells were implanted in the striatum. Both trophic and differentiation factors enhanced the yield of dopamine neurons in the ventral mesencephalic grafts (Torres et al. 2005).

### 2.2.3 MSCs as Cell Therapy for Parkinson's Disease

Mesenchymal stem cells are multipotent stem cells with the capacity for differentiation into various cell types including neurons (MacKenzie & Flake 2002; Choi et al. 2012). These cells can be obtained from human umbilical cord, bone marrow, or adipose tissue (Fu et al. 2002; Kestendjieva et al. 2008; Li et al. 2010; Włodarski et al. 2012). Autotransplantation is possible for cell therapy because the cells can be obtained from the same patient. One interesting immunosuppressive property of these cells is the prevention of an inflammatory response, thus avoiding transplanted cell rejection (Guo et al. 2006; Mathew et al. 2003). Other interesting properties of these cells are their neuroprotective capacity mediated by the delivery of neurotrophic factors (Wilkins et al. 2009; Whone et al. 2012). Several studies have shown that the presence of neurotrophic factors protects the transplanted neurons and favors NSC proliferation and differentiation (Park et al. 2008). These features make MSCs good candidates for cell-replacement therapy in PD. Bone marrow-derived MSCs can be induced to secrete neurotrophic factors such as brain-derived neurotrophic factor (BDNF) and glial cell line-derived neurotrophic factor (GDNF). When these neurotrophic factor-secreting cells were transplanted into the lesioned brain of a Parkinson's disease rat model, regeneration of the damaged dopaminergic nerve terminal in the striatum was observed (Sadan et al. 2009). On the other hand, Li and colleagues differentiated human umbilical vein mesenchymal stem cells (HUVMSCs) into dopaminergic-like cells. Utilizing a rat model of PD, the authors observed that the transplanted cells together with NGF administration considerably increased grafted cell survival (Li et al. 2010).

#### 2.2.4 Neural Stem/Progenitor Cells

The transplantation of fetal or embryonic tissue has significant limitations for use as a cell-replacement therapy such as ethical issues due to the use of human fetal/ embryonic tissue, deleterious immune reactions by the central nervous system after intracerebral transplantation, and the high quantity of cells necessary for an efficient transplantation. In addition, this treatment could have side effects such as dyskinesia and a risk of tumor formation. Consequently, researchers have focused

their efforts on isolating and differentiating in vitro neural stem cells before performing the transplantation. Therefore, NSCs would be obtained first and, later, dopaminergic cells. Thus, fetal brain neural SC-derived dopaminergic cells were observed to have little risk of tumor formation and immune rejection in contrast to stem cells. In addition, neural stem/progenitor cells (NSPCs) are an alternative cell source for neural transplantation because they display important neuroprotective and immunomodulator features that favor survival of the remaining or transplanted neurons (Bonnamain et al. 2012).

Stem cell differentiation generates progenitors of a specific lineage cell such as NSCs. Neural stem cells are multipotent cells with the capacity to proliferate and give rise to cell types with restricted neural lineages (Bai et al. 2007). NSCs differentiate into neural progenitor cells (NPCs) and finally into dopaminergic neurons (Riaz et al. 2002). Although NSCs have a more restricted self-renewal potential, they proliferate in culture forming neurospheres and may be differentiated either before or after transplantation. Both neural stem cell and neural progenitor cells (NSPCs) are present in neurospheres. It has been discussed whether the term NSCs, neural precursor cells, or NPCs are the same cells. However, it is not clear which exact cell types are present within the neurospheres. Recently, it was demonstrated that neurospheres can be maintained in a culture with different cell populations (Narayanan et al. 2012). Currently, it is believed that the cell identities present in neurospheres might differ between species (Steffenhagen et al. 2011).

The NSPCs can be derived from embryonic nerve tissue, adult nerve tissue such as the subventricular zone or carotid body, or from differentiated embryonic/fetal stem cells (Okabe et al. 1996; Pardal et al. 2007; Ara et al. 2010). Autologous transplantation of NSCs derived from adult tissue is an attractive cell therapy against Parkinson's disease (Arias-Carrión and Yuan 2009). The intrastriatal transplantation of NSCs isolated from the developing human CNS in adult rats with unilateral dopaminergic lesions partially ameliorated lesion-induced behavioral deficits in some animals (Svendsen et al. 1997). Pre-differentiated neural progenitor cells are more effective in enhancing the therapeutic potency for PD (Wang et al. 2004). Others studies have shown that transplanted NSCs survive into the substantia nigra pars compacta or striatum in PD rat models (Zhu et al. 2009).

NSCs were isolated from the adult pig subventricular zone, maintaining their capacities for proliferation, self-renewal, and multipotency. Because the pig is immunologically closest to humans, pigs can be an efficient source for cell therapy (Liard et al. 2009). Interestingly, the use of xenografts of porcine neural precursor cell in a rat model of Parkinson's disease showed long-term survival of grafts in the host brain (Harrower et al. 2006).

## 2.2.5 Other Stem Cell Sources

From bone marrow, it is possible to obtain various types of stem cells such as MSCs and hematopoietic stem cells. One study showed that bone marrow-derived stem cells transfected with tyrosine hydroxylase and a GFP-containing plasmid can be later differentiated to neurons. Injection of these cells into the brain of a Parkinson's disease model rat showed favorable results (Zou et al. 2010). On the other hand,

stem cells can be isolated either from the dermis or hair follicles. Both cell types, dermis-derived neural crest-related and epidermal neural crest stem cells, can be differentiated into cells with distinct cell lineages including dopaminergic neuron-like cells. Transplanting these cells into Parkinson's disease model rats showed encouraging results (Kubo et al. 2009). In addition, implantation of epidermal neural crest stem cells into a brain injury rat model leads to cell differentiation into immature astrocytes, which secrete neurotrophic factors (Jackson et al. 2010).

## 2.3 Other Cell Sources for Parkinson's Disease Therapy

Although the tremendous potential of stem cells as an effective cell therapy in Parkinson's disease is well established, cells other than stem cells have also been investigated as alternative sources for cell therapy. These cells may be appropriate for use as supportive or replacement cells for neurons. Among these are amniotic epithelial cells, retinal pigment epithelium (RPE) cells, and fetal dopamine neurons (Sheng et al. 1993; Mendez et al. 2005; Yang et al. 2009; Kakishita et al. 2000). The ability of some cells to secrete neurotrophic factors, differentiate into dopaminesecreting cells in vitro, survive in the striatum in vivo, and increase dopaminergic marker expression levels in Parkinson's disease model rats makes them an excellent candidate for cell therapy against Parkinson's disease (Mendez et al. 2005; Stover and Watts 2008; Yang et al. 2009). The clinical results obtained from the transplant of fetal dopamine neurons in patients with Parkinson's disease were very interesting. When these neurons were transplanted into the striatum or both the striatum and substantia nigra, they presented as functional dopaminergic neurons at the transplantation sites. Likewise, postmortem analyses showed the survival of the transplanted dopaminergic neurons (Mendez et al. 2005).

Despite the progress achieved in this field, numerous complications have prevented the practical application of stem cell-based therapy: ethical problems of fetal and embryonic tissue use, low safety due to teratoma formation risk (Kuroda et al. 2013), and rejection issues and little efficacy caused by very low cell survival. Before the clinical application of cell therapy for patients with Parkinson's disease, safety and efficacy studies, standardized protocols that utilize animal compound-free defined media for culture in vitro, and differentiation procedures that allow us to isolate primarily dopaminergic neurons are required.

# 2.4 Differentiation of Stem Cells into NSCs and Dopaminergic Neurons

It is essential to increase our knowledge on nervous system development and to understand how numerous factors participate in the intrinsic and extrinsic signaling pathways inducing cell differentiation. As a result of these advances, we can design more efficient strategies for cell differentiation in vitro in the future. The differentiation of stem cells involves the formation of neural precursor cells or neural progenitor cells, which can be expanded in vitro. Later, these cells are differentiated further into dopaminergic neurons (Cho et al. 2008). During the differentiation process, neuroepithelial rosette structures are formed (Zhang & Zhang 2010). It was found that not only are stem cells potentially tumorigenic, but neuroepithelial rosettes are as well. An efficient protocol for neuronal differentiation that eliminates ESCs and neuroepithelial rosettes is necessary to avoid risk of tumors or teratoma formation before transplantation. Although some studies performed in rats have shown that multipassage neural precursor cells do not form tumors, low survival and loss of the dopaminergic neuronal phenotype make them unsuitable for replacement therapies (Ko et al. 2009; Jo et al. 2009; Pawitan 2011).

In general, numerous protocols have been described to differentiate stem cells into dopaminergic neurons or dopaminergic-like cells. These protocols have presented with different degrees of efficiency (Hwang et al. 2010). Some protocols utilized a co-culture with PA6 or MS5 mouse stromal cells, and others used chemically defined media or the specific activation of an intrinsic pathway (Momčilović et al. 2012; Watmuff et al. 2012). Currently, there is notable heterogeneity in the yield of tyrosine hydroxylase (TH)-positive cells (Hwang et al. 2010). Although iPSCs and ESCs can be differentiated using the same transcriptional network with same set of morphogens, the differentiation occurs with reduced efficiency and increased variability (Hu et al. 2010). Derivation of only one cell population and increased efficiency during this process are important factors to consider.

Stromal feeder cell-based differentiation protocols have been described by several authors (Perrier et al. 2004; Sonntag et al. 2007). Stromal cells are a source of stromal cell-derived inducing activity (SDIA). Recent work has identified stromal candidate factors inducing the neuronal differentiation of hiPSCs such as hepatocyte growth factor (HGF), stromal cell-derived factor-1 $\alpha$  (SDF1 $\alpha$ ), secreted frizzled-related protein 1 (sFRP1), and vascular endothelial growth factor D (VEGFD) (Schwartz et al. 2012).

Recent evidence has shown that Lmx1a overexpression generates enriched populations of human A9-subtype ventral midbrain DA neurons from hESC or hiPSCs, which should be suitable for cell-replacement strategies (Sánchez-Danés et al. 2012a). Similarly, other studies have shown how overexpression of the nuclear receptor Nurr1 (a transcriptional factor specific to midbrain dopamine (DA) neuronal development) in NPCs isolated from the subventricular zone (SVZ) could induce a DA neuron phenotype (Shim et al. 2007).

As a consequence of the progress in this area, it has been determined that a diverse combination of growth factors and molecules contributes to the expansion and/or differentiation of either stem cells or neural progenitor cells. Among those described include basic fibroblast growth factor (bFGF), fibroblast growth factor 2 (FGF2), fibroblast growth factor 8 (FGF8), fibroblast growth factor 20 (FGF20), docosahexaenoic acid, all-trans-retinoic acid (ATRA), sonic hedgehog (SHH), transcription factor Nato3, a 3 % oxygen level, and ascorbic acid (Volpicelli et al. 2004; Correia et al. 2007; Cooper et al. 2010; Morizane et al. 2010; Chang et al. 2012; Stacpoole et al. 2011; Li et al. 2012; Nissim-Eliraz et al. 2012; Wang et al. 2013).

Similarly, several studies have shown how the bone morphogenic protein antagonist noggin, sonic hedgehog (SHH), FGF8, and Wnt5a promote the differentiation, survival, and functional integration of stem cell-derived DA neurons in vivo (Jensen et al. 2008; Parish et al. 2008; Chiba et al. 2008). Likewise, the BMP, TGF- $\beta$ , and Wnt signaling pathways are involved in midbrain dopamine differentiation. The BMP and TGF- $\beta$  pathway members participate how upstream regulators of the What signaling pathway during midbrain dopamine differentiation of human pluripotent stem cells (Cai et al. 2013). Likewise, another study showed that BDNF contributes to NSC proliferation and differentiation through the activation of the Wnt/ $\beta$ -catenin signaling pathway (Chen et al. 2013). Wnts, a family of secreted proteins, are involved in stem cell differentiation (Prakash and Wurst 2007). It was reported that loss of Wnt1 generates the loss of Lmx1a and Ngn2 expression, affecting the formation of DA neurons in the midbrain floor plate (Andersson et al. 2013). In addition, it has been observed that the Wnt signaling pathway leads to the inhibition of glycogen synthase kinase 3 (GSK-3). GSK-3 inhibition in the presence of epidermal growth factor (EGF) and basic fibroblast growth factor (bFGF) increases hiPSC-NP proliferation. However, an absence of these growth factors generates a switch to neuronal differentiation and a dopaminergic fate (Esfandiari et al. 2012).

The effect of retinoic acid in the neuronal differentiation has been described. It was shown that retinoic acid induced neuronal differentiation of NTERA2 cells through the modulation of Sox2 and Sox3 gene expression. The authors showed downregulation of Sox2 and upregulation of the Sox3 gene (Stevanovic 2003).

Utilizing different stem cell types, it is possible to give rise to dopamine neurons. Recently, Li and colleagues were able to derive dopamine neurons from human cord blood-derived multipotent stem cells (CB-SCs) before treatment with all-trans-retinoic acid (ATRA) (Li et al. 2012). Similarly, the isolation and differentiation of MSCs from the umbilical cord matrix tissue alleviate Parkinsonian symptoms in a rodent model in vivo (Shetty et al. 2013).

On the other hand, iPSCs may also become an important source of dopaminergic neurons. Several differentiation protocols performed in embryonic stem cells were applicable to iPSCs. Thus, it was observed that docosahexaenoic acid stimulated the differentiation of murine iPSCs into dopaminergic neurons (Chang et al. 2012). The neurons obtained from reprogrammed fibroblasts improved symptoms in a rat model of Parkinson's disease (Wernig et al. 2008). Likewise, differentiated Parkinson patient-derived iPSCs reduce motor asymmetry in parkinsonian rats (Hargus et al. 2010).

## 3 Induced Pluripotent Stem Cells (iPSCs)

Although most of our tissues contain adult stem cells that differentiate into any cell type when required, including different tissues (Abdanipour et al. 2011; Darabi et al. 2013), the ability for renewing damaged cells is limited. Over time, the amount of adult stem cells decreases and, thus, the ability to regenerate damaged tissues also decreases. A greater understanding of the molecular mechanisms

and signaling pathways involved in the maintenance and division of these adult stem cells could help design strategies to maintain the tissue regenerative capacity. Certainly, this strategy could be the solution to many diseases involving cell loss. Unfortunately, knowledge in this field remains scarce (Beltrami et al. 2007).

Currently, global research efforts have focused on designing strategies to develop stem cells from different sources. In previous work, different technologies to develop patient-specific stem cells, such as somatic cell nuclear transfer or cell fusion, have been described. In these methodologies, either the somatic nuclear content is transferred to an oocyte or ES cells are fused with somatic cells (Wilmut et al. 1997; Tada et al. 2001; Cowan et al. 2005). Unfortunately, both methods involve technical and ethical considerations that hinder the technology. Likewise, the isolation of adult or embryonic stem cells is not easy to realize, mainly due to technical difficulties and cultural issues. Currently, iPSC technology is a new tool for increasing our understanding of the molecular mechanisms involved in the development of several disorders. Thus, there is great interest in researching the possible use of iPSCs, not just as model of PD but also in future clinical applications (Sun et al. 2010; Stadtfeld and Hochedlinger 2010; Ebben et al. 2011; Han et al. 2011). Currently, these technologies have generated several PD patient-derived iPSC lines (Sánchez-Danés et al. 2012b; Byers et al. 2012). These iPSC lines will certainly help our understanding of the molecular mechanisms of PD and help us to design patient-specific stem cell treatments.

# 3.1 History

In the year 2006, a remarkable finding was reported by Takahashi and Yamanaka in the Yamanaka group in Japan. They showed, for the first time, that an adult somatic cell could be reprogrammed into the pluripotent state by reproducing the ESC signaling pathways (Takahashi and Yamanaka 2006). To define the transcription factors necessary for inducing the reprogramming of somatic cells, the Yamanaka group initially selected a total of 24 separate genes. Previous experience suggested that these genes could play a role in ESC identity development and maintenance (Schöler et al. 1989; Avilion et al. 2003; Rizzino 2009). Thus, they determined a final "cocktail" of transcription factors required to induce pluripotent cell development. A number of these genes are involved in rapid cell proliferation, and some are considered oncogenes. After the analysis of different transcription factor combinations, the authors found that only four factors are essential to reprogram an adult somatic cell to the undifferentiated state, a so-called induced pluripotent stem cell (iPSC). Thus, the authors developed iPSCs by inducing overexpression through the retroviral transduction of Oct4, Sox2, Klf4, and c-Myc in mouse embryonic and adult fibroblasts and subsequent selection for Fbx15 expression (Takahashi and Yamanaka 2006). Shortly after, these studies were successfully confirmed by Wernig and colleagues (Wernig et al. 2007). Subsequently, new selection markers were reviewed for identifying iPSC colonies. The authors showed that selection for Nanog expression generates a germline-competent iPSCs with characteristics similar to ESCs. Under these conditions, they found that the four transgenes (Oct3/4, Sox2, c-Myc, and Klf4) are silenced in Nanog iPSCs (Okita et al. 2007).

Due to the rapid progress in iPSC research, the Yamanaka and Thomson groups reported in 2007 the generation of iPSCs from human cells. These findings are particularly interesting because the contribution of pluripotent cells to disease modeling, drug development, or regenerative medicine was surprising. The Yamanaka group generated iPSCs from adult human dermal fibroblasts using the same four factors: Oct3/4, Sox2, Klf4, and c-Myc (Takahashi et al. 2007). Tumor development is linked with the reactivation of the c-Myc transgene, which is present in the reprogramming cocktail. The presence of this gene could increase tumorigenicity in both the chimeras and progeny mice, which is a serious hurdle for the therapeutic application of iPSCs technology. However, the Thomson group reported that four factors (Oct4, Sox2, Nanog, and Lin28) are sufficient for reprogramming human somatic cells to pluripotent stem cells. Interestingly, Klf4 and c-Myc were not included in the reprogramming cocktail, and the technical limitations that generate viral integration were also eliminated (Yu et al. 2007). Furthermore, new investigations were able to generate human iPSCs from adult dermal fibroblasts without c-Myc (Yu et al. 2007; Nakagawa et al. 2008).

In accordance with the previous descriptions from the Yamanaka, Jaenisch, or Thomson groups, the reprogramming process involved simultaneous overexpression of four transcription factors. In early studies, the authors utilized Oct4, Sox2, Klf4, and c-Myc as pluripotency factors (Takahashi and Yamanaka 2006; Takahashi et al. 2007; Wernig et al. 2007). Thereafter, the Thompson group achieved reprogramming using different pluripotent factors such as Nanog and Lin28 (Yu et al. 2007). Later, reprogramming was also obtained using a fewer number of transcription factors related to stem cell pluripotency (Patel and Yang 2010; Stadtfeld and Hochedlinger 2010; Robinton and Daley 2012). Moreover, some researchers achieved reprogramming including only one or two factors (Grinnell et al. 2007; Kim et al. 2008, 2009a, b, c).

On the other hand, iPSCs have been generated from cells of numerous origins and different species. First, iPSCs were obtained from mice and humans; later, they were generated from other species: rat, pig, rhesus monkey, marmoset, canine, sheep, and snow leopard (Stadtfeld and Hochedlinger 2010; Verma et al. 2012; Sumer et al. 2011; Piedrahita et al. 2011). Certainly, the reprogramming technology offers powerful new opportunities for research not only in the biomedical area. Similarly, iPSCs have been derived from different somatic cell types such as embryonic fibroblasts, adult fibroblasts, hepatocytes, epithelial cells, keratinocytes, melanocytes, mature B cells, mature T cells, pancreatic B cells, and the hepatic endoderm (Nakagawa et al. 2008; Shi et al. 2008; Silva et al. 2008; Stadtfeld et al. 2008a; Guo et al. 2009). Undifferentiated cells have also been useful for reprogramming to iPSCs. These cells include neural stem cells, myeloid progenitor cells, hematopoietic stem cells, adipose-derived stem cells, and cord blood stem cells (Stadtfeld and Hochedlinger 2010). Because the iPS cells can be differentiated into any type of cell with three germ layers, they can be

a cell source relevant for replacement therapy for several disorders. Low transformation efficiency has been a main problem in obtaining these cells.

These findings were startling and altered our knowledge of the epigenetic landscape. Although there has been extensive research to understand how these factors induce changes in gene expression profiles and chromatin modifications through global epigenetic changes, the molecular mechanisms involved in these dedifferentiation processes are incompletely understood (Bernstein et al. 2006).

#### 3.1.1 Oct3/4

The octamer-binding transcription factor POU5F1 (also known as Oct3, Oct4, Oct3/4) is encoded by the POU5F1 gene and expressed during early mouse embryogenesis (Schöler et al. 1989; Palmieri et al. 1994). Oct3/4 has been described as a crucial regulator maintaining pluripotency, which in conjunction with other transcription factors such as Nanog and Sox2, forms a transcription factor molecular network that regulates ESC pluripotency (Boyer et al. 2005; Niwa 2007). Previous studies have shown that Oct3/4 at later stages is expressed in adult stem cells and is considered a pluripotent cell marker (Tai et al. 2005). However, Oct3/4 expression is not limited to stem cells, as differentiated cells also express it (Zangrossi et al. 2007). The differential expression of the isoforms Oct3/4 as Oct3/4A and Oct3/4B may correlate with different processes such as proliferation and differentiation. In fact, a recent study showed that the isoform Oct3/4A is expressed in the nucleus, whereas Oct3/4B is expressed in the cytoplasm in the developing tooth germ (Nakagawa et al. 2012).

Oct3/4 is an essential reprogramming factor, allowing mature somatic cells to return to a pluripotent state as observed in stem cells (Takahashi and Yamanaka 2006; Yu et al. 2007; Nakagawa et al. 2008). Interestingly, their extraordinary ability to reprogram somatic cells into pluripotent cells can be achieved either alone or in collaboration with other factors (Kim et al. 2008, 2009a, b; Li et al. 2011). Currently, a recent study has demonstrated that transcriptional activation of specific genes by Oct3/4 without the gene repression required for differentiation is sufficient for promoting pluripotency (Hammachi et al. 2012). These results contradict those previously reported by researchers who described that the repression of genes required for differentiation is an essential role of Oct3/4 (Boyer et al. 2006; Bilodeau et al. 2009). Oct3/4 does not appear to be required for somatic stem cell self-renewal (Berg and Goodell 2007; Lengner et al. 2007). Moreover, it was observed that Oct3/4 prevents the differentiation and expansion of progenitor cells when overexpressed in adult epithelia (Hochedlinger et al. 2005). Oct3/4 cooperates with a wide set of cofactors. Together with Sox2, Oct3/4 can form a protein-DNA ternary complex that is able to promote the transcriptional activation of fibroblast growth factor 4 (FGF4) (Yuan et al. 1995). Moreover, between Oct4 might promote interaction and Sox2 expression an of itself (Okumura-Nakanishi et al. 2005) and other factors such as Nanog (Kuroda et al. 2005). More recently, Oct3/4 together with Sox2 interacts with a multi-subunit stem cell coactivator complex (SCC) that affects Nanog and Oct4 gene expression as well as genomic regions occupied by Oct4 and Sox2 (Fong et al. 2011).

#### 3.1.2 Sox2

The Sox genes consist of a family of transcriptional regulators involved in the control of nervous system development. Among these is the protein Sox2, an HMG box transcription factor, which is predominantly expressed in undifferentiated cells such as ESCs and is required for the maintaining pluripotency and/or self-renewal (Rizzino 2009).

Studies have shown that Sox2 can form a ternary complex with Oct3 on the FGF4 enhancer DNA sequences promoting FGF4 gene and Nanog gene transcriptional activation. FGF4 is one protein expressed in early embryonic development (Yuan et al. 1995; Kuroda et al. 2005). Likewise, a Sox2 regulatory region, which is also under the control of this Oct3/4-Sox2 complex, was identified (Tomioka et al. 2002). It is believed that the Oct3/4 and Sox2 expression levels are maintained by the Oct4/Sox2 complex in pluripotent cells by a reciprocal transcriptional regulation mechanism (Chew et al. 2005). It has been proposed that the major role of Sox2 is to regulate multiple transcription factors that affect Oct3/4 expression, thus maintaining Oct3/4 expression in pluripotent cells (Masui et al. 2007). Importantly, the transcription factor Sox2 and Oct3/4 have been implicated in early lineage decisions. While both factors cooperate to suppress trophoblast fates in ESCs, they also establish the three lineages present at implantation (Avilion et al. 2003; Li et al. 2007). Sox2 is highly expressed in dividing neural progenitor cells throughout the central nervous system (CNS) (Ellis et al. 2004; Ferri et al. 2004; Komitova and Eriksson 2004). Thus, Sox2 is expressed in embryonic early neural precursors of the ventricular zone and in the ependymal of adult cells. Sox2 is also expressed in lower proportions in differentiated neurons. An important role for Sox2 in neuron maintenance has been described (Ferri et al. 2004). Indeed, Sox2 expression is characteristic of NSCs in the adult brain; however, not all NSCs have the capability to differentiate into all neural cell types in vivo (Brazel et al. 2005). An interesting difference in Sox2 gene expression levels between human and mouse has been described. Mouse primordial germ cells (PGCs) express Oct3/4, Nanog, and Sox2, whereas human PGCs express Oct3/4 and Nanog but not Sox2 (Perrett et al. 2008; De Jong et al. 2008). Sox2 is considered a neural stem cell marker for ICM stem cells. Sox2 has been used as a key transcription factor for reprogramming somatic cells into induced pluripotent cells, but it was not considered essential (Kim et al. 2008, 2009a). Reprogramming neural progenitor cells into iPSCs can be achieved by the combination of factors necessary for reprogramming. Depending on the cellular context, exogenous Sox2 expression is not necessary if cells already express high levels of Sox2 (Eminli et al. 2008). Likewise, Sox2 is not absolutely necessary for neural stem cell self-renewal or multipotency (Miyagi et al. 2008). On the other hand, Sox2 has also been shown to be implicated in tumorigenesis in various organs (Lengerke et al. 2011). These observations indicate that there is a relationship between pluripotency and tumor development (Bernhardt et al. 2012).

### 3.1.3 Nanog

The transcription factor Nanog plays a key role during lineage specification and in the maintenance of pluripotency in both ICM and ESCs. Nanog in conjunction with other pluripotent factors such as Oct3/4 and Sox2 controls expression of target genes that

possess important functions in pluripotency development. Thus, it was found that Nanog-deficient ESCs lost their ability to self-renew (Cavaleri and Schöler 2003; Mitsui et al. 2003). High levels of Nanog can allow for ESC self-renewal and prevent differentiation in the absence of cytokine LIF or feeder layers (Pan and Thomson 2007). These key transcription factors form a complex regulatory network affecting the expression levels of different target genes related to pluripotency or differentiation (Wang et al. 2006). It was observed that Oct3/4 and Sox2 binding to the Nanog promoter is essential for Nanog transcription in mouse and human embryonic stem cells (Rodda et al. 2005). Moreover, it was observed that Nanog regulates downstream genes necessary for both pluripotency and the differentiation of stem cells through two distinct transactivation domains (Pan and Pei 2003). Other studies described that Nanog dimerization is necessary for interaction with several proteins that promote stem cell pluripotency (Wang et al. 2008). In fact, Nanog is a transcriptional activator for Rex-1, which is a marker for undifferentiated ESCs (Shi et al. 2006). When embryonic stem cells are differentiated into primitive endoderm, Nanog is repressed. Thus, Nanog downregulation plays a crucial role in cell fate during the early embryonic development (Hamazaki et al. 2004). Several studies have shown that Nanog expression is not exclusive to pluripotent cells because it is also present in tumor cells and plays a novel role in tumor development (Jeter et al. 2009). While mouse Nanog expression is high in undifferentiated ESCs in the ICM of the blastocyst, germ cell tumor, and teratoma-derived cell lines, expression is low in several adult tissues (Hart et al. 2004; Hoei-Hansen et al. 2005). Thus, Nanog levels may act as a control mechanism to decide between cell self-renewal and differentiation. Moreover, the expression of pluripotent genes such as Nanog in both seminomas and breast carcinomas suggests a direct role in the progression of different types of carcinomas and could serve as an important marker of tumorigenesis (Ezeh et al. 2005). It was observed that many characteristics present in stem cells are also present in cancer cells. This suggests that any changes in the complex regulation mechanism existing in pluripotent cells could lead to tumor development. Furthermore, it was found that Nanog priming before full reprogramming may generate germ cell tumors (Grad et al. 2011). It is fundamental to discriminate both cell types correctly. On the other hand, regulation of the pluripotent state involves the participation of various factors. Thus, many endogenous noncoding RNA molecules, polycomb complexes, and DNA methylation may play a role in regulating the pluripotent state (Kashyap et al. 2009). Tay and colleagues described how different microRNAs targeting Nanog, Oct4, and Sox2 coding regions participate in embryonic stem cell differentiation (Tay et al. 2008). Likewise, the identification of conserved lncRNA-encoding genes such as lncRNAs, AK028326, and AK141205 was described as direct targets of Oct4 and Nanog (Sheik Mohamed et al. 2010).

## 3.2 Delivery Systems into Donor Cell

Certainly, the main problem in generating iPS cells is an extremely inefficient transformation process. Thus, iPSCs reprogramming is influenced by numerous

variables that must be considered before choosing a reprogramming method. In fact, reproducibility and the quality of the resulting iPSCs are also affected by the delivery system.

Currently, there are several strategies to generate iPS cells, as rapid progress has been made in this field. These methods utilize several mechanisms for factor delivery as non-integrating vectors, excisable vectors, retrovirus, adenovirus, microRNA, and even soluble proteins, and each has different reprogramming efficiencies (Okita et al. 2008; Stadtfeld et al. 2008b; Kaji et al. 2009; Soldner et al. 2009; Woltjen et al. 2009; Zhou and Freed 2009). Several systems have been described to deliver transgenes into donor cells including integrative and non-integrative delivery systems. Integrative methods have been described such as viral delivery vectors, lentiviral delivery vectors, transfection of linear DNA, and the piggyBac transposon (González et al. 2011; Robinton and Daley 2012; Bellin et al. 2012). Wide variations in the efficiency of reprogramming were observed if reprogramming was performed in cells at different differentiation states. Less differentiated cells as progenitor cells can be reprogrammed more easily. Moreover, less differentiated cells may require fewer transcription factors to achieve reprogramming (Stadtfeld and Hochedlinger 2010).

The firsts iPSCs were obtained by retroviral transduction using pMX-based retroviral vectors (Takahashi and Yamanaka 2006; Takahashi et al. 2007; Okita et al. 2007; Nakagawa et al. 2008). In this case, the retroviruses expressing each gene separately showed an iPSC generation efficiency of less than 0.02 % when either mouse or human fibroblasts were reprogrammed (Stadtfeld and Hochedlinger 2010). Likewise, polycistronic vectors were designed resulting in the simultaneous, almost stoichiometric, expression of transcription factors encoded by the polycistron.

Later, other groups utilized effective lentiviral vectors as delivery systems of reprogramming factors (Yu et al. 2007). In contrast to other retroviruses, lentiviral vectors have a higher infection efficiency, allowing for the infection of both dividing and nondividing cells. Moreover, the development of new lentiviral vectors such as Tet-inducible reprogramming lentiviruses or the STEMCCA lentiviral vector permits the successful ectopic expression of reprogramming factors (Major et al. 2011; Sommer et al. 2012). However, there are several limitations to the safe use of lentiviral vectors for reprogramming. Viral integration into the genome can be random, thus affecting expression of tumor suppressor genes or oncogenes. Therefore, the possibility that viral transgenes can ultimately be spontaneously reactivated increases the risk of tumor formation (Okita et al. 2007).

To avoid viral integration, new alternatives have emerged for factor delivery through a non-integrative method. Adenoviral reprogramming of mouse fibroblasts or liver cells to iPSCs has been described (Stadtfeld et al. 2008b; Okita et al. 2008). Likewise, human iPS cells were derived using a non-integrating episomal vector. Because of the removal of the episoma, the human iPSCs were free of vector and transgene sequences (Yu et al. 2009; Chou et al. 2011). Human iPSCs were also derived using non-viral minicircle DNA vectors (Okita et al. 2010). The major

advantage of these technologies is the lack of genomic integration. One of the main issues in obtaining iPSCs is not just the lower reprogramming efficiency, but also choosing a method that does not change the genomic DNA. Other interesting reprogramming methods involve the use of proteins (Zhou et al. 2009). Although the technique has been described as having a very low reprogramming efficiency, multiple hiPSC lines have been derived through protein-based reprogramming. These cells rescued motor deficits in a PD model (Rhee et al. 2011).

One of the methods that have been proposed to significantly increase reprogramming efficiency is the use of RNA and certain molecules (Warren et al. 2010; Wang et al. 2012).

# 4 Direct Reprogramming into Neurons

One of the problems with the use of induced pluripotent stem cells is that they can generate tumors. Recently, different groups of researchers have shown that only a few transcription factors were sufficient to generate the direct conversion of fibroblasts to functional neurons without generating a pluripotent stem cell stage (Kim et al. 2011). These cells were less likely to form tumors, avoiding much of the carcinogenic risks of pluripotent cells; they are called induced neuronal (iN) cells. Vierbuchen and colleagues analyzed a pool of 19 neural-lineage-specific transcription factors as possible candidates for direct reprogramming. They showed that the combination of three factors, Ascl1, Brn2, and Myt11, is able to transform mouse embryonic and postnatal fibroblasts into functional neurons in vitro. The iN cells generated expressed neuronal markers such as GABA (Vierbuchen et al. 2010). They were also capable of converting fetal and postnatal human fibroblasts into iN cells in the presence of the basic helix-loop-helix transcription factor NeuroD1 (Pang et al. 2011). Functional neurons were also obtained from adult human fibroblasts (Pfisterer et al. 2011). Similarly, Caiazzo and collaborates identified three other transcription factors, Mash1, Nurr1, and Lmx1a, that also can be utilized to induce the reprogramming of mouse and human fibroblasts into dopaminergic neurons (Caiazzo et al. 2011). On the other hand, another group of investigators found that the combination of five transcriptional factors, Mash1, Ngn2, Sox2, Nurr1, and Pitx3, were capable of reprogramming human fibroblasts into DA neuron-like cells (Liu et al. 2012).

One of the major disadvantages of this technology in cell-replacement therapy is the low number of cells generated. A new procedure that directly reprograms mouse fibroblasts into neural progenitor cells could avoid this problem. Unlike iN cells, these cells can be expanded in vitro (Kim et al. 2011).

Furthermore, before being used for cell transplantation, the cells must survive at the transplantation site in addition to being effective in the animal model. However, this tool is an attractive option for other cell types for generating patient- and disease-specific neurons to be used in a cell disease models.

## 5 Conclusion

In recent decades, research efforts have focused on replacing dopaminergic neurons that are lost in patients with Parkinson's disease. To this end, cellular therapies that replace degenerated cells have been the major focus and may be promising therapeutic options in patients with Parkinson's disease in the future. Preliminary studies have utilized adult tissue, embryonic/fetal tissue, and embryonic stem cells as a source of dopaminergic cells. However, safety issues such as the risk of tumor formation and immunological rejection have prevented this type of technique from being an effective treatment against Parkinson's disease. Advances in iPSC research have allowed scientists to obtain specific cells from the same patient, avoiding rejection problems. However, obstacles remain for its clinical use such as the lack of standardized protocols that result in a dopaminergic cell phenotype that grafts without undifferentiated cells and without a risk of tumor formation. Direct reprogramming of somatic cells into induced neuronal (iN) cells without generating a pluripotent stage could be a solution that avoids tumor risk. However, as a result of the low quantity of cells obtained, more investigations are necessary to turn this technique into an efficient cell-replacement therapy. If we overcome these obstacles, iPS and iN cells may be, without a doubt, a very promising therapy in the near future.

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# Alpha Synuclein in Parkinson's Disease

Christine Lund Kragh, Marina Romero-Ramos, Glenda Halliday, and Poul Henning Jensen

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#### Abstract

The perception of Parkinson's disease (PD) as a disease centered on dopaminergic striatonigral neurodegeneration has changed fundamentally since 1997 when the first mutation in the SNCA gene (PARK1) encoding  $\alpha$ -synuclein was discovered (Polymeropoulos et al. 1997). This discovery formed the basis for a new description of brain pathology characterized by the presence of  $\alpha$ -synuclein

C.L. Kragh (🖂) • M. Romero-Ramos • P.H. Jensen

G. Halliday

Department of Biomedicine, Aarhus University, Aarhus C, Denmark e-mail: clp@biokemi.au.dk; mrr@biokemi.au.dk; PHJ@BIOKEMI.AU.DK

Neuroscience Research Australia and University of New South Wales, Sydney, NSW, Australia e-mail: g.halliday@neura.edu.au

aggregates in brain cell inclusions that are the hallmarks of PD and other synucleinopathies: dementia with Lewy bodies (DLB) and multiple system atrophy (MSA). This field has been thoroughly covered by many reviews during the last decade (Gai et al. 1998; Spillantini and Goedert 2000; Huang et al. 2004; Ubhi et al. 2011). This review will briefly highlight the historical breakthroughs but focus on  $\alpha$ -synuclein modifications, human neuropathology, biomarker potential, current animal models and the new concepts emerging after the significance of extracellular  $\alpha$ -synuclein has gained support.

#### Keywords

Animal models • Biomarkers • Dementia with Lewy bodies • Lewy bodies • Multiple systems atrophy • Parkinson disease • Phosphorylation • Truncation •  $\alpha$ -synuclein

## 1 Introduction

The perception of Parkinson's disease (PD) as a disease centered on dopaminergic striatonigral neurodegeneration has changed fundamentally since 1997 when the first mutation in the SNCA gene (PARK1) encoding  $\alpha$ -synuclein was discovered (Polymeropoulos et al. 1997). This discovery formed the basis for a new description of brain pathology characterized by the presence of  $\alpha$ -synuclein aggregates in brain cell inclusions that are the hallmarks of PD and other synucleinopathies: dementia with Lewy bodies (DLB) and multiple system atrophy (MSA). This field has been thoroughly covered by many reviews during the last decade (Gai et al. 1998; Spillantini and Goedert 2000; Huang et al. 2004; Ubhi et al. 2011). This review will briefly highlight the historical breakthroughs but focus on recent developments in concepts and hypotheses.

## 2 Genetics

Missense mutations in the SNCA gene substituting single amino acids (A30P, E46K, and A53T) cause early-onset autosomal dominant PD and DLB (Polymeropoulos et al. 1997; Kruger et al. 1998; Zarranz et al. 2004). Additionally, genomic multiplications of the SNCA gene have been found in patients with early-onset PD and DLB (Singleton et al. 2003) demonstrating that simple overexpression and perhaps developmentally dysregulated expression of the normal protein is sufficient to initiate disease development. The discovery of these rare cases was essential because they causally tie  $\alpha$ -synuclein to the development of a PD syndrome, leading to the widely accepted hypothesis that a gain of function in  $\alpha$ -synuclein contributes to the pathogenesis of PD. A possible role for  $\alpha$ -synuclein in sporadic PD is corroborated by a series of genome-wide association studies carried out on ethnically diverse populations of PD patients. The single-nucleotide

polymorphisms conferring the strongest risk are in the SNCA gene (Simon-Sanchez et al. 2009). Although SNCA is the strongest risk gene, polymorphisms within this gene only increase the risk 2–3 times above the low prevalence of about 0.14 % but serve to corroborate a mechanistic link between the development of sporadic PD and  $\alpha$ -synuclein.

# **3** Structure of α-Synuclein

 $\alpha$ -Synuclein is a small 140-amino acid protein, which was first identified as associated with purified synaptic vesicles from the electric ray *Torpedo californica* (Maroteaux et al. 1988). It is one of three synuclein family members ( $\alpha$ ,  $\beta$ , and  $\gamma$ ) identified in humans. The  $\alpha$ -synuclein protein can be divided into two regions: (i) the N-terminal region (residues 1–95), which contains seven imperfect repeats with a highly conserved hexameric motif (KTKEGV) and the aggregation-prone non-amyloid component (NAC) sequence (residues 60–95), and (ii) the unstructured C-terminal region (96–140), which is rich in acidic residues (Ueda et al. 1993; George et al. 1995).

The heat-stable nature of human  $\alpha$ -synuclein was already described in 1994 (Jakes et al. 1994), and it was ascribed to its natively unfolded structure that possesses an ability to form  $\alpha$ -helical structures in apolar solvents (Weinreb et al. 1996) and upon interactions with acidic phospholipid membranes (Clayton and George 1998). When analyzed by gel filtration, the unfolded structure caused the 14.4 kDa  $\alpha$ -synuclein peptide to elute corresponding to an approximately 55 kDa globular protein. Moreover, certain dimers appearing on denaturing SDS-polyacrylamide gels were due to misincorporation of a cysteine residue in some preparations of bacterially expressed human  $\alpha$ -synuclein (Masuda et al. 2006). Despite its unfolded nature, purified  $\alpha$ -synuclein has been demonstrated to exhibit dynamic long-range interactions between the C-terminus and the central NAC domain and even further toward the N-terminus (Cho et al. 2009). Moreover, it undergoes a dramatic structural change upon binding to acidic liposomes where the N-terminal repeat regions fold into two  $\alpha$ -helices (Bussell and Eliezer 2003; Chandra et al. 2003).

 $\alpha$ -Synuclein aggregates into amyloid fibrils by a nucleation-dependent mechanism and the disease-causing mutations as well as C-terminal truncations favor the consumption of monomers into aggregates. The transformation from a monomeric state to an amyloid state proceeds through soluble oligomeric intermediates (Wood et al. 1999). Several lines of evidence suggest that the oligomers are the most toxic species. First, the disease-causing A30P mutation accelerates the initial oligomerization of  $\alpha$ -synuclein in vitro, whereas it retards the formation of mature filamentous aggregates (Conway et al. 2000). Second,  $\alpha$ -synuclein oligomers bind synthetic vesicles and cause transient permeabilization representing a potential mechanism of neurotoxicity (Lashuel et al. 2002). This was corroborated when oligomeric forms of  $\alpha$ -synuclein applied extracellularly were shown to trigger

calcium entry and toxicity in a neuronal cell line (Danzer et al. 2007). Third, cytotoxicity in cellular models is usually seen without heavily aggregated  $\alpha$ -synuclein (Xu et al. 2002; Outeiro et al. 2008; Tetzlaff et al. 2008; Kragh et al. 2009). The protofibrils formed in vitro from purified  $\alpha$ -synuclein represent a diverse group generated from simple aggregation of monomers in buffer to induction by various agents and conditions, e.g., apolar solvents, lyophilization, and oxidative cross-linking by, e.g., metals and oxidation products from lipid peroxidation (Sharon et al. 2001; Nasstrom et al. 2011). Immunological characterization of  $\alpha$ -synuclein oligomers has been feasible using two different antibodies: the pan-oligomeric A11 antibody recognizing soluble precursors of many types of amyloid-type fibrils (Kayed et al. 2007) and the FILA-1 antibody recognizing  $\alpha$ -synuclein-specific epitopes shared between filaments and soluble oligometers. The FILA-1 antibody recognizes elevated levels of soluble  $\alpha$ -synuclein aggregates in DLB brain extracts (Paleologou et al. 2009) indicating commonalities between in vitro-formed oligomers and those present in vivo. At present, the structural relation between in vitro-formed  $\alpha$ -synuclein protofibrils and those found in cells and brain tissue is unclear and structure-specific imaging agents are in high demand.

Two recent reports challenge the dogma of  $\alpha$ -synuclein as a natively unfolded protein and show evidence supporting native tetramers rich in  $\alpha$ -helical structure. Both reports indicate that  $\alpha$ -synuclein from red blood cells and recombinantly expressed in *Escherichia coli* with a short N-terminal extension can be purified as folded tetramers that irreversibly dissociate into monomers (Bartels et al. 2011; Wang et al. 2011). Interestingly, the tetramers are unable to aggregate and amyloid-type aggregation is only possible after dissociation into monomers. Importantly, the PD-causing missense mutations destabilize the tetramers and thus favor aggregation. These findings are controversial but potentially important as they hypothesize that a stabilization of the native tetrameric state could be a fruitful therapeutic strategy.

# 4 Posttranslational Modifications

 $\alpha$ -Synuclein is subject to different posttranslational modifications. It can be phosphorylated at Ser87, Ser129, and Tyr125 (Ellis et al. 2001; Anderson et al. 2006; Paleologou et al. 2010) in human brain tissue. Though predominantly found in a non-phosphorylated state in vivo,  $\alpha$ -synuclein is phosphorylated at Ser129 in inclusions in all  $\alpha$ -synucleinopathies (Fujiwara et al. 2002; Anderson et al. 2006). Both Ser129-phosphorylation and ubiquitination are highly enriched in  $\alpha$ -synuclein aggregates in diseased brain, whereas other less abundant modifications occur in both soluble and insoluble fractions.

The role of Ser129-phosphorylation in promoting toxic  $\alpha$ -synuclein aggregation is unclear as in vitro phosphorylation of this residue has been demonstrated to prevent amyloid-type insoluble aggregation (Fujiwara et al. 2002;

Paleologou et al. 2008). α-Synuclein phosphorylation at Ser129 accelerates inclusion formation and/or toxicity in cell culture models (Smith et al. 2005b; Sugeno et al. 2008; Kragh et al. 2009) and correlates with pathology in transgenic Drosophila (Chen and Feany 2005). Studies in rats expressing the phosphomimic form of  $\alpha$ -synuclein (S129D) using adeno-associated virus (AAV) have yielded conflicting results (see section on "Animal Models"). However, S129D may not be a suitable phosphomimicking variant of  $\alpha$ -synuclein as it was demonstrated that S129D does not reproduce the effects of phosphorylation on structural and aggregation properties of  $\alpha$ -synuclein in vitro (Paleologou et al. 2008). Mechanistically, Ser129-P may contribute functionally by conferring novel ligand-binding properties as demonstrated using a phosphopeptide co-immunoprecipitation approach (McFarland et al. 2008). Phosphorylation of tyrosines also occurs in  $\alpha$ -synuclein (Ellis et al. 2001) and Tyr125-phosphorylation decreases with age in human brain tissue (Chen et al. 2009). Deletion of the tyrosine residues in  $\alpha$ -synuclein increased toxicity in a Drosophila model (Chen et al. 2009) indicating a protective role for tyrosine phosphorylation that may be lost with aging. To fully understand the consequence of  $\alpha$ -synuclein phosphorylation, it is important to identify the involved kinase(s). Casein kinase II, GRK2/5, and polo-like kinases 2/3 have been shown to phosphorylate  $\alpha$ -synuclein in vitro, in cell culture, *Drosophila*, and mice (Pronin et al. 2000; Chen and Feany 2005; Arawaka et al. 2006; Inglis et al. 2009; Mbefo et al. 2010) (for a review see Oueslati et al. 2010).

Proteolytic C-terminal truncations after residues 115, 119, 122, 133, and 135 have been demonstrated in human brain tissue (Anderson et al. 2006), and such modifications increase  $\alpha$ -synuclein aggregation in vitro (Crowther et al. 1998). Interestingly,  $\alpha$ -synuclein is a substrate for cytoplasmic calpains, and cleaved  $\alpha$ -synuclein is found in PD and DLB brain (Mishizen-Eberz et al. 2005; Kim et al. 2006; Dufty et al. 2007), suggesting that calpain may generate truncated toxic species of  $\alpha$ -synuclein.

The early focus on dopaminergic neurons in substantia nigra in PD prior to the description of the Braak hypothesis and the use of L-dopa for therapy forms the basis for a large series of experiments on dopamine effects on  $\alpha$ -synuclein and vice versa (for a review Leong et al. 2009). Key findings are that dopamine stimulates the formation of soluble off-pathway  $\alpha$ -synuclein oligomers in vitro that do not aggregate into fibrils (Conway et al. 2001) and the dopamine oxidation product dopaminochrome promotes the formation of spherical oligomers (Norris et al. 2005). These observations were corroborated in cellular experiments where the dopamine level was modulated by expression of tyrosine hydroxylase (TH) forms (Mazzulli et al. 2006). In addition,  $\alpha$ -synuclein has been related to different key proteins of the dopaminergic system.  $\alpha$ -Synuclein influences dopamine synthesis by decreasing TH and/or aromatic amino acid decarboxylase activity via changes in the phosphorylation status of these two enzymes (Perez et al. 2002; Tehranian et al. 2006; Lou et al. 2010). Mutated  $\alpha$ -synuclein A53T leads to low levels of VMAT2 and consequently to accumulation of dopamine in neurons (Lotharius et al. 2002).  $\alpha$ -Synuclein levels have also been implicated in regulating the activity of the

dopamine transporter (DAT), although controversies exist. Hence, direct binding between transgenic h $\alpha$ -synuclein and DAT in cells enhanced dopamine uptake and dopamine-dependent cell death (Lee et al. 2001). However, a later study coexpressing the two proteins identified a negative effect on dopamine uptake (Wersinger and Sidhu 2003). Finally, using the opposite approach by silencing endogenously expressed  $\alpha$ -synuclein showed a decreased sensitivity to the DATdependent toxin MPP+ and decreased dopamine uptake (Fountaine and Wade-Martins 2007). Hence, the definitive role of dopamine-dependent  $\alpha$ -synuclein modifications in vivo and the effect of  $\alpha$ -synuclein levels and modifications on dopamine metabolism are still enigmatic.

# 5 Cellular Metabolism

 $\alpha$ -Synuclein is synthesized as a cytosolic protein but it can exist in various subcompartments, e.g., membrane associated or encapsulated with various posttranslational modifications. Natively unmodified  $\alpha$ -synuclein is readily degraded by the 20S proteasome (Liu et al. 2003) and by chaperone-mediated autophagy in lysosomes in vitro (Cuervo et al. 2004). How  $\alpha$ -synuclein is degraded in the brain is less clear, and studies using cell lines and primary neuronal cultures have yielded different results (reviewed in (Xilouri and Stefanis 2011)). Neurons in culture appear to degrade the bulk of  $\alpha$ -synuclein by chaperone-mediated autophagy and macroautophagy (Vogiatzi et al. 2008). A recent study using transgenic mice expressing human  $\alpha$ -synuclein or  $\alpha$ -synuclein-eGFP analyzed the levels of  $\alpha$ -synuclein after topical application of inhibitors of the 20S proteasome and the autophagy-lysosome pathway (Ebrahimi-Fakhari et al. 2011). Using cranial window access to the brain surface combined with live imaging and tissue extraction, it was elegantly demonstrated that the basal  $\alpha$ -synuclein turnover is mediated by the proteasome system and this process appears even more important with aging. Upon stress mediated by increased  $\alpha$ -synuclein expression or compromised proteasomal activity, the autophagy-lysosome pathway is activated (Ebrahimi-Fakhari et al. 2011). The importance of lysosomal dysfunction in relation to Lewy body (LB) formation has been corroborated by the presence of LB pathology in Gaucher disease that is caused by dysfunctional lysosomal hydrolases (Vitner et al. 2010). Exocytosis of vesicle-bound  $\alpha$ -synuclein was demonstrated in 2005 (Lee et al. 2005), and direct neuron-neuron transfer of  $\alpha$ -synuclein was later confirmed using cell transplantation (Desplats et al. 2009). This formed the important experimental frame for testing whether  $\alpha$ -synuclein by a prion-like mechanism causes the neurodegenerative progression described by the Braak hypothesis (Braak et al. 2004) ranging from development of LBs in fetal human dopaminergic neurons transplanted into PD patients (Kordower et al. 2008; Li et al. 2008) to Lewy-like pathology ascending to the brainstem after oral toxin-based generation of  $\alpha$ -synuclein pathology in the gut (Pan-Montojo et al. 2010) and to neuronal uptake of preformed  $\alpha$ -synuclein aggregates in vitro (Volpicelli-Daley et al. 2011).

#### 6 Normal Functions

The normal function of  $\alpha$ -synuclein has been difficult to pinpoint although its effective axonal transport (Jensen et al. 1999) and selective localization in nerve terminals (Maroteaux et al. 1988) suggest a function in this structure. Although it is expressed at high levels in the brain, it is also found in other tissues, e.g., hematopoietic cells (Miller et al. 2004). Reduction of  $\alpha$ -synuclein in cultured neurons reduced the distal pool of synaptic vesicles (Murphy et al. 2000), and knockout mice corroborated a presynaptic function as evidenced by reduced striatal dopamine content and an increased neurotransmitter release upon paired stimuli (Abeliovich et al. 2000). However, the phenotype was surprisingly mild with the most striking feature being a resistance to the toxin MPTP (Dauer et al. 2002). Although knockout of  $\alpha$ -synuclein alone or  $\alpha$ -synuclein and  $\beta$ -synuclein together has minimal effects on neuronal function (Chandra et al. 2004), triple knockout mice lacking all synuclein members demonstrate a robust phenotype with gross reduction in presynaptic size and premature death (Greten-Harrison et al. 2010). These mice display deficits in assembly of SNARE complexes that are required for fusion of synaptic vesicles to the plasma membrane. Interestingly, it was recently shown that the synucleins act as chaperones for presynaptic SNARE proteins (Burre et al. 2010).

Transgenic expression of  $\alpha$ -synuclein has been considered a rational approach for most studies of pathological  $\alpha$ -synuclein effects because of the dominant inheritance of PD and DLB associated to mutations in the SNCA gene. Hence, wild-type and mutated human  $\alpha$ -synuclein has been expressed in a range of cell lines, primary brain cells, yeast cells, and organisms ranging from *Caenorhabditis elegans* and *Drosophila* to mice, rats, and monkeys with variable results. Initially the criteria for success were generation of a LB-like structure and this frustrated many researchers. End points have later changed toward neuronal degeneration and cell death because it was realized that LBs may take months to years to develop in human neurons.

## 7 Pathogenic Functions

Mechanisms by which abnormal accumulation of  $\alpha$ -synuclein disrupts cellular functions have been investigated intensively in several different model systems. Yeast studies have demonstrated that  $\alpha$ -synuclein accumulation leads to a blockade of vesicular transport from the endoplasmatic reticulum to the Golgi apparatus and hereby inhibits the viability of the cells (Cooper et al. 2006). Moreover, activation of an unfolded protein response has been demonstrated in SHSY5Y cells overexpressing  $\alpha$ -synuclein (Sugeno et al. 2008) and in postmortem brain tissue from PD patients (Hoozemans et al. 2007). However, it became clear in the yeast study that the toxic response was dependent on the  $\alpha$ -synuclein dose as revealed by comparing high and low expressers (Su et al. 2010). A different approach has been taken in mammalian cells that tolerate expression of  $\alpha$ -synuclein but where cytotoxicity was induced by coexpressing proteins known to stimulate  $\alpha$ -synuclein aggregation like p25 $\alpha$  and synphilin-1 (Engelender et al. 1999; Kragh et al. 2009). Using this approach it was demonstrated that p25 $\alpha$  induced cellular degeneration that could be attenuated by inhibitors of  $\alpha$ -synuclein aggregation as well as of Ser129-phosphorylation (Kragh et al. 2009). The approach of including inhibitors of  $\alpha$ -synuclein aggregation may be readily applicable to other models and thereby allow phenotypes caused by mere overexpression of  $\alpha$ -synuclein to be more clearly distinguished from those forming aggregations.

It has previously been difficult to identify soluble  $\alpha$ -synuclein aggregates in cellular models. Nevertheless, they have been detected by Western blotting in mesencephalic neurons exposed to polyunsaturated fatty acids (Sharon et al. 2003) and in a dopaminergic cell line overexpressing A53T  $\alpha$ -synuclein (Zhou and Freed 2005). Moreover, a fluorescence lifetime imaging-based technique to probe the organization of  $\alpha$ -synuclein within the cell has demonstrated the formation of small oligomers (Klucken et al. 2006). Toxic effects have also been demonstrated by applying  $\alpha$ -synuclein from culture media to cells, but the toxic pathways elicited by internal versus external  $\alpha$ -synuclein are still unclear. An elegant approach was used to test whether mature aggregates or insoluble protofibrils are responsible for toxic effects caused by overexpression of  $\alpha$ -synuclein. In this study,  $\alpha$ -synuclein mutants designed to favor generation of soluble protofibrils exhibited increased toxic potential compared to wild-type  $\alpha$ -synuclein in cell and animal models (Karpinar et al. 2009).

Several lines of evidence implicate a dysfunction of the ubiquitin-proteasome system in the pathogenesis of  $\alpha$ -synucleinopathies. Studies show that LBs contain ubiquitinated proteins (Chung et al. 2001; McNaught and Jenner 2001) and that the proteasomal activity is decreased in the substantia nigra of PD patients (McNaught and Jenner 2001; McNaught et al. 2003). Moreover, aggregated  $\alpha$ -synuclein has been demonstrated to inhibit the activity of the proteasome in vitro (Snyder et al. 2003; Lindersson et al. 2004; Emmanouilidou et al. 2010), and this may have significant effects on proteasomal-dependent degradation of short-lived proteins.

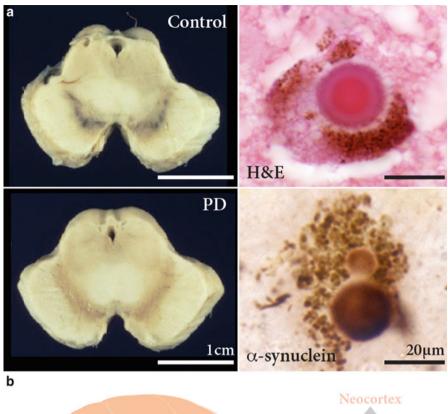
Mitochondrial dysfunction as a consequence of  $\alpha$ -synuclein accumulation has also been a major area of interest. Abnormalities of mitochondrial function and increased free radical-mediated damage were described in PD brain before the first PD-associated gene mutations were discovered.  $\alpha$ -Synuclein can localize to mitochondria (Hsu et al. 2000; Tanaka et al. 2001; Smith et al. 2005a) and  $\alpha$ -synuclein expression increases cellular sensitivity to rotenone, a mitochondrial complex I inhibitor (Orth et al. 2003; Ved et al. 2005).  $\alpha$ -Synuclein has also been shown to reduce ATP synthesis and mitochondrial membrane potential (Kamp et al. 2010). Moreover, structural abnormalities of mitochondria have been observed in transgenic mice overexpressing mutant  $\alpha$ -synuclein (Martin et al. 2006).

The existence of cellular  $\alpha$ -synuclein excretion and uptake by neighboring neurons as previously described may form a path for spreading of the disease process in the nervous system. The potential of such mechanisms has been corroborated by the dramatic neuronal spread of  $\alpha$ -synuclein pathology in transgenic mice expressing human A53T mutant  $\alpha$ -synuclein (Luk et al. 2012). Unilateral injection of homogenates from old symptomatic M83 mice into the brains of young mice caused a rapid propagation of aggregated and hyperphosphorylated  $\alpha$ -synuclein ranging from the olfactory bulb to the spinal cord that also traversed the corpus callosum to the contralateral hemisphere. The propagation required the recruitment of endogenous mouse  $\alpha$ -synuclein. Surprisingly, similar pathology was induced upon injection of in vitro-formed insoluble aggregates of a fusion protein consisting of C-terminally truncated human  $\alpha$ -synuclein (1–120) and a Myc epitope tag that lacked the S129 phosphorylation site. Hence, insoluble  $\alpha$ -synuclein aggregates may possess "prion-like" seeding activity (Luk et al. 2012).

## 8 Human Pathology

PD is diagnosed by two separate abnormalities in the absence of sufficient neuropathology for an alternative neurodegenerative disorder (Dickson et al. 2009). A moderate to substantial loss of the dopaminergic neurons in the substantia nigra is required, a finding that also occurs in the majority of movement disorders, and the presence of hallmark neuronal LBs in brainstem predilection sites is also required (Fig. 1a). LBs are also the hallmark pathology of DLB with the two conditions separated mainly on clinical rather than pathological grounds, particularly at end stage (McKeith et al. 2005). PD patients have a dominant slowly progressive movement disorder with cognitive impairment often intervening very late in the disease when LBs infiltrate into the cortex (Dickson et al. 2009). DLB patients have a dominant more rapidly progressive dementia disorder and have cortical LBs often with coexisting Alzheimer's disease (Halliday et al. 2011a). Although  $\alpha$ -synuclein forms abnormal aggregates in all synucleinopathies, neuronal LBs only occur in PD and DLB with  $\alpha$ -synuclein inclusions in alternate cell types (oligodendroglia) or cellular locations (axonal spheroids) in other synucleinopathies (Halliday et al. 2011a). The reason for the different locations and of neurodegeneration involving  $\alpha$ -synuclein remain mechanisms largely unexplored.

In addition to PD and DLB, LBs occur in a small proportion of asymptomatic elderly, a finding which has allowed the development of a hypothesized pathological staging scheme for PD (Braak et al. 2003) and one for both PD and DLB (Beach et al. 2009). Braak hypothesis for the progression of pathology in PD states that the pathology starts peripherally in the olfactory and autonomic nervous systems and progresses through connected regions into the spinal cord and medulla oblongata (stage 1), progressing over time to the pons (stage 2) and midbrain (stage 3), to the basal ganglia (stage 4) and limbic systems (stage 5), and finally to association cortices (stage 6), largely sparing the somatosensory and motor system (Braak et al. 2003) (Fig. 1b). The alternate hypothesis for the progression of LB pathology across PD and DLB by Beach and colleagues (Beach et al. 2009) states that pathology begins in the olfactory bulb (stage 1) and then progresses in two different directions depending on clinical phenotype – for the motor phenotype (PD) LBs





Mesocortex, allocortex

# Midbrain dopamine forebrain, diencephalon

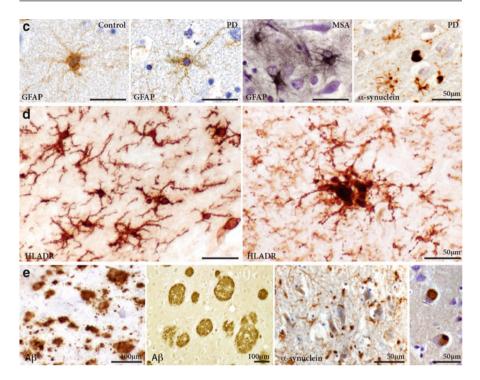
Pontine tegementum

# Medulla oblongata

Peripheral and enteric nervous system

700

Fig. 1 (continued)



**Fig. 1** Neuropathology found in patients with  $\alpha$ -synucleinopathies. (a) The two diagnostic pathologies required for PD are the loss of pigmented dopaminergic neurons in the midbrain substantia nigra (shown macroscopically at left) and LB formation in remaining brainstem pigmented neurons (shown at right). Macroscopic photos of transverse sections through the midbrain from a control and a patient with PD demonstrate the loss of the dark neuromelanin pigment in the ventral midbrain. LBs within remaining pigmented neurons in patients with PD can be visualized as large round intracellular inclusions observed with hematoxylin and eosin (H&E) or  $\alpha$ -synuclein immunohistochemistry. (b) Diagrammatic representation of the Braak staging scheme for the progression of PD from the olfactory and peripheral nervous systems to invade the brain in a slow hierarchical fashion from the lower brainstem to the upper brainstem and forebrain to the cortex. (c) Photomicrographs of the typical astrocytic changes that occur in PD versus MSA. While reactive astrogliosis is typical in MSA (observed as an increase in GFAP and hypertrophy of their cell bodies compared to controls), this reaction is not typically observed in end-stage cases of PD. Rather, in PD a large number of astrocytes abnormally accumulate  $\alpha$ -synuclein throughout their cytoplasm, a finding associated with the formation of the large round intracytoplasmic LBs in nearby neurons. (d) Photomicrographs of the microglial changes observed in PD. Amoebic, phagocytic microglia aggregate only in regions with neuronal loss like substantia nigra (*right*), while in areas abnormally accumulating  $\alpha$ -synuclein, the microglia display more HLA-DR reactivity in association with an increased branching morphology (left). (e) Pathologies associated with each other in patients with the more rapidly progressing DLB and later onset PD with early dementia. The density of amyloid plaques in both the striatum (*left*) and cortex (middle left) is related to each other and to the severity of dementia in these patients. Lewy neurites in the hippocampus (middle right) are also related to the severity of plaque and LB pathology (at right) in such cases

infiltrate the brainstem (stage 2a) and for the dementia phenotype (DLB) LBs infiltrate the limbic system (stage 2b). Finally, for both phenotypes, LBs are found in both the brainstem and limbic system (stage 3) prior to infiltrating cortical regions (stage 4) (Beach et al. 2009).

The Braak staging of PD pathology has been controversial due to discrepancies in a number of validation studies and also due to some conceptual difficulties, while the alternate staging scheme has not been independently evaluated. However, it should be noted that the majority of longitudinally followed cases with motordominant PD fit the Braak staging [see review (Halliday and McCann 2010)]. The most obvious difficulty with the Braak staging scheme is using asymptomatic cases with LBs to only stage the progression of PD and not DLB. The more restricted number of stages in the alternate hypothesis incorporating DLB makes this staging scheme of limited current utility. It should be noted that in  $\sim 10$  % of asymptomatic elderly, LBs occur only in cortical regions (Zaccai et al. 2008). Additionally, the clinical progression of PD relates to the severity of dopaminergic cell loss (Greffard et al. 2006). This may not be surprising if LBs mark neurons destined to die with all LB debris removed following cell death. However, if this was the case, then as the disease progresses, there should be a reduction of LB pathology as cell death is enhanced and affects more regions, rather than the concept of pathology progressing to accumulate in more and more neurons and cells still viable in the brain. Empirical data shows that even in regions thought to be heavily affected with pathology, only 4-5 % of neurons contain LBs in PD (Harding et al. 2002; Greffard et al. 2010). Furthermore, there is no widespread brain tissue loss in PD (Weintraub et al. 2011), and the pattern and severity of cell loss does not reflect the demonstrated pattern and severity of LB formation. Definitive proof of the progression of LB pathology will only occur with the development of biomarkers capable of identifying this pathology.

In addition to the cell loss and LB pathology required for diagnosis, glial changes are obvious in PD (Halliday and Stevens 2011). Astrocytes in PD disengage from neurons leaving the neuronal membranes more exposed (Knott et al. 1999). They undergo PD-specific changes in that they accumulate non-fibrillar  $\alpha$ -synuclein and lack the typical reactive astrogliosis that usually accompanies neurodegeneration (Song et al. 2009) (Fig. 1c). In particular, there is often no obvious hypertrophy of astrocytes or upregulation of their glial fibrillary acidic protein (GFAP) or any substantial proliferation, a finding that differs significantly from reactive astrogliosis observed in MSA (Song et al. 2009) (Fig. 1c). This may suggest that any reactive trigger resolves or is removed over time in PD, possibly by cellular uptake. In contrast, the astrocytes in patients with MSA do not accumulate  $\alpha$ -synuclein but have the typical reactivity observed in chronic neurodegenerative conditions with obvious cell hypertrophy and upregulation of GFAP (Song et al. 2009) (Fig. 1c), changes that are associated with long-lasting reorganization of tissue architecture. The accumulation of  $\alpha$ -synuclein in astrocytes in PD is related to the severity of LB infiltration and synaptic degeneration (Braak et al. 2007),

affecting more astrocytes ( $\sim$ 45 %; (Song et al. 2009)) compared with neurons ( $\sim$ 4–5 %; Harding et al. 2002; Greffard et al. 2010).

 $\alpha$ -Synuclein in astrocytes causes an upregulation of inflammatory mediators (Klegeris et al. 2006; Lee et al. 2010) as well as neuroprotective factors (Power et al. 2002; Ishida et al. 2006; Durrenberger et al. 2009; Michael et al. 2011). The increase in  $\alpha$ -synuclein deposition also correlates with an increase in activation of microglia (Croisier et al. 2005), and with the removal of astroglial processes from neurons, the vacated perineuronal space becomes occupied by amoeboid phagocytic microglia (Knott et al. 1999) (Fig. 1d). While activated microglia are well described as a feature of PD (see review Hirsch et al. 2012), phagocytic microglia are only found in regions of cell loss and express scavenger and high-affinity IgG receptor Fc $\gamma$  (Orr et al. 2005) as well as macrophage markers (Croisier et al. 2005). These data suggest that the relatively large number of astrocytes affected by  $\alpha$ -synuclein plays a significant role in the disease process.

PD is one of the age-associated neurodegenerative conditions and occurs often in association with other age-related neuropathologies (Compta et al. 2011). In particular, the pathology most often associated with age-related dementia is amyloid plaque accumulation (Matthews et al. 2009), which also contributes to dementia in older cases of PD (Compta et al. 2011) (Fig. 1e). It should be noted that patients with dementia and sufficient neurofibrillary tangle pathology for a diagnosis of Alzheimer's disease are not considered to have a LB syndrome that contributes to their dementia, due to their substantially greater Alzheimer-type pathology (Dickson et al. 2009). However, amyloid plaque pathology is known to occur in nearly all patients with DLB, being much less frequent in younger PD patients, even those with dementia (Halliday et al. 2011b). This confirms in vivo PiB imaging studies in patients with PD (Gomperts et al. 2008) and suggests that, in such typical patients, the  $\alpha$ -synuclein cortical LB pathology is sufficient for their late end-stage dementia. Therefore, there is considerable clinical heterogeneity in patients with LBs that associates with other age-related pathologies – patients with a younger onset often have a slow disease course and more pure LB pathology at end stage, while patients with an older onset are more likely to have additional age-related pathologies and a more rapid and complex clinical course, the most rapid course occurring in patients with DLB (Compta et al. 2011).

### 9 α-Synuclein in Human Fluids as a Biomarker for PD Progression

Aggregated  $\alpha$ -synuclein accumulates in degenerating neurons in PD making measurements of  $\alpha$ -synuclein levels a rational biomarker for diagnosis and prediction of disease susceptibility.  $\alpha$ -Synuclein has been detected in blood, CSF, and saliva at low nanogram/ml levels (e.g., (Fjorback et al. 2007; Devic et al. 2011; Shi et al. 2011)). Most investigations have focused on detecting the total amount of  $\alpha$ -synuclein but have also investigated the presence of polymers using identical monoclonal anti- $\alpha$ -synuclein antibodies as catching and detecting antibody (Tokuda et al. 2010). Future developments will likely aim at detecting specific  $\alpha$ -synuclein isoforms, e.g., misfolded species as detected in brain extracts, and specific phosphorylations to allow for better stratification. For a review on  $\alpha$ -synuclein in CSF, please see (Mollenhauer et al. 2010).

## 10 Modeling PD

Different organisms from flies and worms to rodents and primates have been used to model PD based on expression of wild-type or mutated human  $\alpha$ -synuclein. In this section, we will highlight the discoveries related to mammals and refer readers to a recent review about other approaches (Pienaar et al. 2010). There are numerous human  $\alpha$ -synuclein transgenic mice although the associated phenotypes are not always consistent probably due to the varying levels and areas of transgene expression as a consequence of the promoters used (for further reading see (Magen and Chesselet 2010; Chesselet and Richter 2011)). Disease modeling has focused on the following points: neurodegeneration in general and dopaminergic neurodegeneration in particular, basal ganglia-associated motor behavior and signs of abnormal  $\alpha$ -synuclein folding and handling (see Table 1 for overview).

#### 10.1 Dopaminergic Neurodegeneration

Mice lines using the dopaminergic TH promoter have been rather disappointing in modeling PD (Matsuoka et al. 2001; Rathke-Hartlieb et al. 2001) and only doublemutated  $\alpha$ -synuclein (A30P and A53T; a condition not found in humans) has been able to induce dopaminergic failure (Richfield et al. 2002), and this was even obtained with the pan-neuronal Thy1 promoter (Ono et al. 2009). Interestingly, expressing C-terminal-truncated  $\alpha$ -synuclein (1–120 or 1–130  $\alpha$ -synuclein) under the TH promoter induced dopaminergic cell death, although not progressively and related to developmental defects (Tofaris et al. 2006; Wakamatsu et al. 2007; Wakamatsu et al. 2008). Using the rat AAV model, the ability of truncated  $\alpha$ -synuclein to promote pathological accumulation of wild-type  $\alpha$ -synuclein was corroborated (Ulusoy et al. 2010). More widely expressing promoters like Thy1 and PDGF have also been used to model dopaminergic degeneration and subtle phenotypes have been obtained (Masliah et al. 2000; Rockenstein et al. 2002; Lam et al. 2011).

An alternative to the classic transgenic mice lines is adult transgenesis using viral vectors to induce overexpression of wild-type or mutated forms of  $\alpha$ -synuclein in nigrostriatal neurons. This leads to progressive and selective neurodegeneration and dopaminergic loss in the basal ganglia (Kirik et al. 2002; Lo Bianco et al. 2002; Yamada et al. 2004; Maingay et al. 2006). These models have shown characteristics of  $\alpha$ -synuclein misfolding (Lo Bianco et al. 2004;

Promoter	Human α-syn	a-Syn in SN	Dopaminergic system	Pathological α-syn	Behavioral phenotype	Remarks	References
Rat-TH	120 Trunc	+	Striatal TH terminal loss	Filamentous	Motor	Microgliosis	Tofaris et al. (2006)
			¢DA	Insoluble (WB)			
	130 Trunc	+	Nigral and striatal TH neuronal loss	ż	Motor		Wakamatsu et al. (2008)
	A30P or A53T	+	No change	ż			Matsuoka et al. (2001),
			↑MPTP				Manning-Bog et al.
			susceptibility in				(2003), Yu et al. (2008)
			ICCA and				
			↓ Paraquat				
			susceptionity				
	A30P and A53T	+	Nigral and striatal Ubiquitin (-) Motor TH neuronal loss	Ubiquitin (–)	Motor	Microgliosis preceding Richfield et al. (2002), cell death Su et al. (2009)	Richfield et al. (2002), Su et al. (2009)
			†Susceptibility to	Congo red		Long-lasting increase	
			MPTP or maned	- (-)		on proinflammatory	
			and paraquat			cytokines	
	A30P	+	No change	ż			Rathke-Hartlieb et al.
			Unchanged MPTP suscentibility				(2001)

Table 1	Table 1 (continued)						
Promoter	Human α-syn	a-Syn in SN	Dopaminergic system	Pathological α-syn	Behavioral phenotype	Remarks	References
Mouse prion	E46K	+		Lewy body- like	Motor correlated with spinal cord-brainstem	Microgliosis	Emmer et al. (2011)
				Fibrillar	pathology		
				PSer129			
				a-syn			
				Insoluble			
				(WB)			
				Nitrated			
				Tau inclusions			
	A30P	SN?	↓ Capacity of the	ż	Abnormal motor	Changes in NE	Yavich et al. (2004, 2005,
			DA storage pool		response to L-dopa	neurotransmission	2006), Oksman et al.
			and evoked release				(2009)
		Str+	No change in TH				
			staining in Str or SN				
	A53T	+	ż	Not	Motor	Lamp 1+ (if null	Gispert et al. (2003),
				aggregated		background)	Cabin et al. (2005)
	A53T		↑ D1 receptor in	Ubiquitin (+)	Hyperactivity associated	Degenerated	Lee et al. (2002),
			SN	ThioF S (+)	with DA	mitochondria	Martin et al. $(2006)$ ,
				Insoluble			Unger et al. (2006)
				(WB) 0.:			
				Uligomeric			
				Truncated	Motor defect associated \UDACOMPLEX IV	↓Complex IV	
			↓ DA reuptake and Nitrated	Nitrated	with spinal cord-		
			DAT in Str	Spheroids	brainstem	Altered expression on	
				Eosinophilic	nemonegeneration	inflammation-related genes	
						0	

Giasson et al. (2002), von Coelln et al. (2006), Norris et al. (2007), Tsika	et al. (2010), Gao et al. (2011)		Zhou et al. (2008)					Rockenstein et al. (2002),	Fleming et al. (2004),	Fleming and Chesselet	(2006), Fernagut et al. (2007), Fleming et al.	(2008)			(continued)
If crossed with DJ1 KO no change	Parkin KO no change	†Susceptibility (maned + paraquat)						Microgliosis		Early	electrophysiological Changes in Str			I	
Severe associated to motor neurons			Motor and cognitive					Sensorimotor				Motor	Olfactory	Colonic motility	
Oligomeric and insoluble (WB)	Oxidized Filamentous Nitrated	Regional different oligomeric	Ubiquitin (+)	PSer129 ¤-svn	Oligomeric	soluble and	insoluble (WB)	Prot.	K resistant	PSer129	a-syn	Oligomeric	(WB)	Insoluble (WB)	
No Change	DAergic hypersensitivity to LPS-induced	inflammation	No change					↑DA release at 6	months			18 moths $\downarrow$ DA	in Str		
+			+					-/+							
A53T			Y39C					ΜT							
			Thy1												

Promoter	Human α-syn	SN	Dopanniergic	r autorogicat α-syn	Behavioral phenotype	Remarks	References
	A30P	+	Normal DA	Prot K racistant	Motor and cognitive	Oxidation of metabolic	Kahle et al. (2001), Neumann et al. (2002)
			vuinvin al all ages	IN IUSISIAIII		uitochondria	Poon et al (2005)
				Insoluble M-svn			Freichel et al. (2007),
				PSer129			Schell et al. (2009)
				α-syn			
				ThioF S (+)			
				Ubiquitin (+)			
	A30P and	+	Nigral and striatal	PSer 129	Motor		Ono et al. (2009)
	A53T		TH neuronal loss	α-syn			
PDGF	WT	-/+	Striatal TH	Non-fibrillar	Motor	↓Neurodegenerative	Masliah et al. (2000),
			terminal loss	a-syn		changes upon	Masliah et al. (2005),
				Oxidized		immunization (active	Koob et al. (2010),
				a-syn		or passive)	Haggerty et al. (2011)
			¢DA	Hyperphosp. TAU			
				Ubiquitin (+)			
Hamster	A30P	+	No changes	Oligomeric	Motor	Microgliosis	Gomez-Isla et al. (2003),
prion			↑ Susceptibility to MPTP	and truncated soluble (WB)			Nieto et al. (2006)

Eslamboli et al. 2007), markers of dysfunctional synapses, and neuroinflammation (Theodore et al. 2008; Chung et al. 2009; Sanchez-Guajardo et al. 2010). So far, this is the only approach that has allowed transgenesis in nonhuman primates to successfully model several of the cardinal symptoms of the disease (Eslamboli et al. 2007; Yasuda et al. 2007).

#### 10.2 Non-dopaminergic Neurodegeneration and Braak

The Braak hypothesis posits that  $\alpha$ -synuclein degeneration also occurs in nondopaminergic neurons (Fig. 1b). Indeed enteric nervous system anomalies and colonic problems have been demonstrated in  $\alpha$ -synuclein mice (Wang et al. 2008; Kuo et al. 2010). Olfactory problems being early symptoms in the disease have also been observed early in mice along with concomitant  $\alpha$ -synuclein pathology in the olfactory bulb (Fleming et al. 2004, 2008; Wang et al. 2008). Very pronounced brainstem and spinal cord degeneration has been observed in the lines using the prion promoter (Giasson et al. 2002; Lee et al. 2002) with up to 75 % of motor neuron cell death (Gispert et al. 2003).

#### **10.3** Pathological α-Synuclein In Vivo

Several  $\alpha$ -synuclein transgenic mice models have shown ubiquitin-positive intracellular inclusions (Masliah et al. 2000; Kahle et al. 2001; Lee et al. 2002; Eslamboli et al. 2007; Zhou et al. 2008) and characteristics of aggregation, like resistance to proteinase K treatment (Magen and Chesselet 2010). Early dysfunctional or plastic changes have been observed in the tissue before pathological deposition of  $\alpha$ -synuclein can be detected that appears to correlate over time with the severity of neurodegeneration (Gispert et al. 2003). The putative role of  $\alpha$ -synuclein Ser129-phosphorylation has been seen in different models (Magen and Chesselet 2010). Phosphorylation of Ser129 is a prerequisite for dopaminergic cell death in *Drosophila* (Chen and Feany 2005), but this has not been corroborated in mammalian models. Three studies have been conducted comparing the neurotoxicity of expressing wild-type and pseudo-phosphorylated S129D mutant human  $\alpha$ -synuclein. In two studies wild-type  $\alpha$ -synuclein was more toxic than S129D, while the third reported no difference between the two  $\alpha$ -synuclein species (Gorbatyuk et al. 2008; Azeredo da Silveira et al. 2009; McFarland et al. 2009). However, it should be kept in mind that S129D  $\alpha$ -synuclein structurally is a poor mimic of Ser129-phosphorylated  $\alpha$ -synuclein (Paleologou et al. 2008). Indirect approaches for increasing Ser129-phosphorylation using overexpression of phosphoprotein phosphatase 2A activity in a Thy1 transgenic mice line (Lee et al. 2011) or the  $\alpha$ -synuclein directed kinase GRK6 using viral vectors in rats (Sato et al. 2011) report increased toxicity, but these approaches also affect the phosphorylation status of other proteins. Noteworthy, phosphorylation at Ser129 is not necessary for  $\alpha$ -synuclein

transgene-dependent cell death to occur, as demonstrated in animals overexpressing  $\alpha$ -synuclein C-terminally truncated before the Ser129 residue (Wakamatsu et al. 2008; Ulusoy et al. 2010).

#### 10.4 Inflammation and Microglia in the Disease

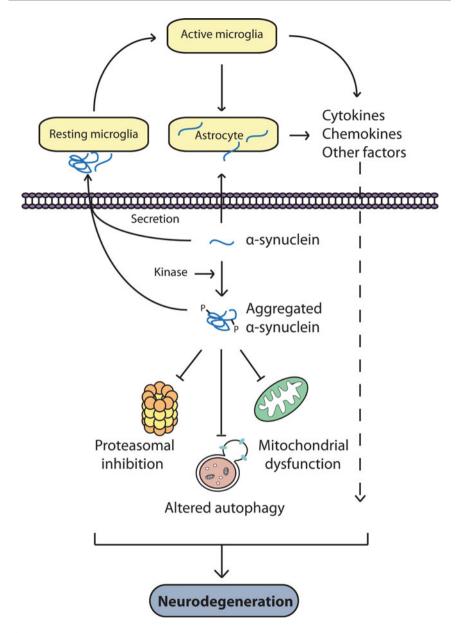
Microglia activation and neuroinflammation in PD is recapitulated in several models: changes in microglia numbers or morphology, the cytokine pattern, or the gene expression (van der Putten et al. 2000; Neumann et al. 2002; Gomez-Isla et al. 2003; Tofaris et al. 2006; Su et al. 2008; Emmer et al. 2011). The microglial activation patterns correlate with the degree of  $\alpha$ -synuclein-dependent neuropathology in the rat viral vector PD model (Sanchez-Guajardo et al. 2010), and prodegenerative synergism exists between the proinflammatory environment induced by local or peripheral LPS treatment and  $\alpha$ -synuclein in transgenic models (Gao et al. 2008; Gao et al. 2011). Neuroprotection observed by both passive and active vaccination in an  $\alpha$ -synuclein transgenic mice line demonstrates that the peripheral immune system may be harnessed therapeutically and such an approach is already in a Phase 1 trial (Masliah et al. 2005; Haggerty et al. 2011).

#### 10.5 Interactions with Other Brain Disease-Related Genes

Regarding the possible interactions with other proteins and genes associated with PD, coexpression of A53T  $\alpha$ -synuclein and LRRK2 (wild-type, mutated G2019S, or KD) resulted in synergistic toxicity enhancing and accelerating the  $\alpha$ -synuclein-induced neurodegeneration (Lin et al. 2009). However, no difference in toxicity was observed in A53T  $\alpha$ -synuclein transgenic mice when crossed with DJ1-deficient (Ramsey et al. 2010) or parkin null mice (von Coelln et al. 2006). Transgenic lines coexpressing wild-type  $\alpha$ -synuclein and  $\beta$ -amyloid peptides showed more severe anomalies than single transgenics, suggesting a shared pathogenic event between both proteins (Masliah et al. 2001). In parallel, the role of  $\alpha$ -synuclein in Huntington's disease has also been highlighted by a decrease in neurodegenerative changes in a knock-in mouse model of Huntington's disease with an  $\alpha$ -synuclein knockout background (Tomas-Zapico et al. 2011).

### 11 Conclusion

There is still considerable doubt over the species of  $\alpha$ -synuclein that contributes most to neurotoxicity (Fig. 2). Currently there is a huge gap between data obtained through in vitro versus in vivo methods. This is highlighted by recent data that suggests the native state of  $\alpha$ -synuclein is a folded tetramer (Bartels et al. 2011; Wang et al. 2011). If this concept proves true, it has fundamental



**Fig. 2** Proposed mechanisms of neurodegeneration in PD. There are multiple potential pathways of neurodegeneration in PD.  $\alpha$ -Synuclein aggregation and accumulation (e.g., of phosphorylated  $\alpha$ -synuclein) can cause proteasomal, lysosomal, and mitochondrial impairment leading to neurodegeneration. In addition,  $\alpha$ -synuclein can be secreted from neurons and taken up by microglia and astrocytes. Proinflammatory mediators such as cytokines and chemokines produced by activated microglia activate astrocytes, and the products released by activated microglia and astrocytes may exert neurotoxic effects

implications for conceptualizing the potential molecular mechanisms involved in  $\alpha$ -synucleinopathies. How Ser129-phosphorylation contributes to these structures or to pathogenesis needs to be determined, as Ser129-phosphorylation is tightly linked to human pathological lesions. To date the functional significance of Ser129-phosphorylation in cell and in vivo models has been inconclusive, findings that may be related to its structural state. Development of novel structure-specific imaging tools, like antibodies and small molecules, is likely to assist with understanding of both the structural and posttranslational changes in  $\alpha$ -synuclein that occur in  $\alpha$ -synucleinopathies. It is important to clarify the significance of Ser129-phosphorylation of different  $\alpha$ -synuclein structural species to determine their role in disease initiation and progression because kinases hold great potential as drug targets.

The novel hypothesis that the disease is spread through the uptake of extracellular  $\alpha$ -synuclein has significant implications currently driving considerable research activity. The observation that astrocytes seem to be more prone to such uptake in PD now needs to be understood fully. Particularly, the functional significance of the early and pervasive astrocytic accumulation of  $\alpha$ -synuclein needs to be determined. Does such behavior play a paralytic role in the neurodegeneration in PD? Regarding uptake of extracellular  $\alpha$ -synuclein, research questions currently pursued include the following: How is a-synuclein externalized during aging and disease and in which forms? How does external  $\alpha$ -synuclein affect cells at the cell surface and upon internalization? The answers to these questions will be important for the further development of the currently promising in vivo  $\alpha$ -synuclein immunotherapies, as externalized  $\alpha$ -synuclein is more amenable to such therapies. Whether there are specific  $\alpha$ -synuclein species and/or cell types that are optimal for antibody-dependent targeting needs to be determined. Furthermore, any immunotherapy successes require replication in multiple models.

Aging is known to be the greatest risk factor for PD and LB pathology, but the mechanisms affected by aging still need to be identified. While the importance of mitochondrial function with aging needs to be considered, the specificity that mitochondrial aging precipitates only  $\alpha$ -synucleinopathies and not other neurodegenerative conditions requires further explanation. Although speculative, Tyr125-phosphorylation may represent a  $\alpha$ -synuclein-dependent cytoprotective mechanism associated with tetramer stability that is lost upon aging. The identification of the toxic species of  $\alpha$ -synuclein and changes with aging will be important to determine. The identification that  $\alpha$ -synuclein can be catabolized through autophagy and recent in vivo modeling suggests that disruption in the cross talk between proteasomal and autophagic  $\alpha$ -synuclein catabolism may be important, particularly as the catabolism of  $\alpha$ -synuclein changes during brain development and aging. Another significant factor may be the coexisting pathologies that accumulate with aging. Cellular defenses against misfolded proteins have been characterized in model organisms, and their role in PD needs to be clarified. Whether concentrating on  $\alpha$ -synuclein therapeutic interventions alone is the best strategy needs to be clarified.

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# Autophagic Pathways and Parkinson Disease

# Miquel Vila and Marta Martínez-Vicente

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#### M. Vila

Neurodegenerative Diseases Research Group, Vall d'Hebron Research Institute-CIBERNED, Barcelona, Spain

Department of Biochemistry and Molecular Biology, Autonomous University of Barcelona (UAB), Barcelona, Spain

Catalan Institution for Research and Advanced Studies (ICREA), Barcelona, Spain e-mail: miquel.vila@vhir.org

M. Martínez-Vicente (🖂)

Neurodegenerative Diseases Research Group, Vall d'Hebron Research Institute-CIBERNED, Barcelona, Spain

Department of Biochemistry and Molecular Biology, Autonomous University of Barcelona (UAB), Barcelona, Spain e-mail: marta.martinez@vhir.org; martamar9@gmail.com

#### Abstract

Autophagy is responsible for the degradation and recycling of intracellular material including organelles, cytosolic proteins, and accumulated misfolded proteins. Increasing evidence implicates autophagy dysfunction in several neurodegenerative disorders including Parkinson's disease (PD). In this chapter, we analyze recent studies that provide new links between genes associated to PD and the autophagy process. Mutations in PD-associated proteins like  $\alpha$ -synuclein, parkin, PINK1, LRRK2, DJ-1, UCH-L1, ATP12A3, and GBA have been recently directly or indirectly linked to alterations in the autophagic system, thereby contributing to the understanding of the role of this system in the pathogenesis of Parkinson's disease.

#### **Keywords**

α-synuclein • ATP12A3 • Autophagy • DJ-1 • GBA • LRRK2 • Parkin • Parkinson's disease • PINK1 • UCH-L1

### List of Abbreviations

AD	Alzheimer's disease
ALS	Amyotrophic lateral sclerosis
CMA	Chaperone-mediated autophagy
GBase	Glucocerebrosidase protein
GD	Gaucher's disease
HD	Huntington's disease
LB	Lewy bodies
LIR	LC3-interacting region
LMP	Lysosomal membrane permeabilization
LSD	Lysosomal storage disorders
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine
MTOC	Microtubule-organizing center
PD	Parkinson's disease
ROS	Reactive oxygen species
SNpc	Substantia nigra pars compacta
UBA domain	Ubiquitin-associated domain
UPS	Ubiquitin-proteasome system
WT	Wild type

#### 1 Introduction

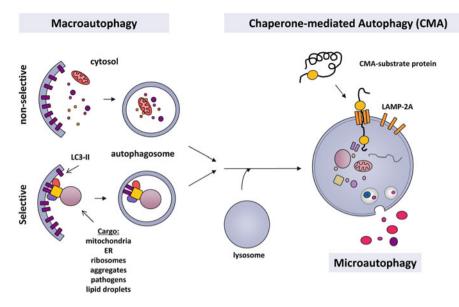
In the last few years, autophagy malfunction had been profusely linked to a growing number of human disorders (Levine and Kroemer 2008; Mizushima et al. 2008), including several neurodegenerative diseases such as Parkinson's disease (Crews et al. 2010; Dagda et al. 2009; Levine and Kroemer 2008; Narendra et al. 2008; Pan et al. 2008a; Stefanis et al. 2001; Vives-Bauza et al. 2009; Webb et al. 2003; Dehay et al. 2010; Vila et al. 2011), Alzheimer's disease (AD) (Boland et al. 2008; Ling and Salvaterra 2009; Nixon et al. 2005; Pickford et al. 2008), Huntington's disease (HD) (Kegel et al. 2000; Nagata et al. 2004; Petersen et al. 2001; Ravikumar et al. 2004), and amyotrophic lateral sclerosis (ALS) (Morimoto et al. 2007).

The first evidence that the lysosomal system could be altered in neurodegenerative disorders came from the observed accumulation of autophagosomes in brain samples from patients and animal models of these diseases (Anglade et al. 1997; Kegel et al. 2000; Nixon et al. 2005; Sapp et al. 1997; Dehay et al. 2010). This increase in autophagosomes was originally attributed to an activation of the autophagic process aimed at eliminating the abnormal accumulation of proteins occurring in these diseases. However, now we know that accumulation of autophagosomes can also be secondary to a failure in the completion of the autophagic process. After some years of abundant research in the field, a better understanding of the role of autophagy in the intracellular quality control system, particularly in neurons, is now providing new evidences to identify the specific autophagic pathways and steps affected in each of the neurodegenerative pathologies. All the accumulated information may provide new therapeutic approaches aimed at modulating autophagy to attenuate, halt, or prevent neurodegeneration.

## 2 Types of Autophagy

Autophagy, from the Greek "auto" (self) and "phagy" (eating), refers to the delivery of intracellular components to lysosomes for degradation. Lysosomes are intracellular organelles which contain acid hydrolases able to break up all type of intracellular and extracellular components and macromolecules (Bohley and Seglen 1992; Dice 2000; Mullins and Bonifacino 2001). In mammalian cells, there are various types of autophagy (Fig. 1) - macroautophagy, microautophagy, and chaperone-mediated autophagy (CMA) – differing in their mechanisms, their specificity of the substrate to be degraded, their regulation, and their function (Cuervo 2006; He and Klionsky 2006; Klionsky 2005; Levine and Kroemer 2008). Although each type of autophagy displays particular features, they all share some common components and are highly interconnected, resulting in a cross talk between the different autophagic mechanisms as well as between other non-lysosomal proteolytic systems, such as the ubiquitin-proteasome system (UPS). This connection between the different proteolytic systems in the cell allows the activation of compensatory mechanisms, when necessary (Kaushik et al. 2008; Massey et al. 2006a; Nedelsky et al. 2008; Pandey et al. 2007).

In *chaperone-mediated autophagy* (CMA), specific cytosolic proteins are directly translocated into the lysosomes through the lysosomal membrane without any vesicular trafficking (Cuervo and Dice 1996; Majeski and Dice 2004; Massey et al. 2004) (Fig. 1). All CMA substrates contain a sequence biochemically related to the pentapeptide KFERQ (Dice 1990), which targets the substrate to the



**Fig. 1** *Autophagy pathways.* Three types of autophagy occur in mammalian cells. In chaperonemediated autophagy (CMA), specific cytosolic proteins containing the CMA-targeting motif are recognized by a chaperone complex and sent to lysosome where they bind the membrane receptor LAMP-2A and directly are translocated into the lysosomes for degradation. Microautophagy involves the sequestration and degradation of complete regions of the cytosol (including proteins and organelles) through invaginations and tubulations of the lysosomal membrane. In macroautophagy intracellular components are enclosed in a double-membrane vesicle called autophagosome. This vesicle is then fused with lysosomes, which provides the hydrolytic enzymes required for the total degradation of the sequestered material. Macroautophagy can be a nonselective process engulfing in-bulk portions of the cytosol but also can be a selective mechanism where specific substrates are recognized by different autophagic adapters and sent selectively to degradation

lysosome membrane with the help of some cytosolic chaperones and binding to receptor membrane protein LAMP-2A (Agarraberes and Dice 2001; Cuervo and Dice 1996). The substrate protein is then translocated into the lysosomal matrix where it is rapidly degraded by the hydrolytic enzymes. CMA is activated by nutrient stress (Cuervo et al. 1995) but also in response to some other stress conditions like oxidative stress or exposure to toxic compounds (Kiffin et al. 2004).

*Microautophagy* involves the sequestration and degradation of complete regions of the cytosol by the lysosomal system, including cytosolic proteins and organelles, through the engulfment of the lysosomal membrane itself (Ahlberg and Glaumann 1985; Marzella et al. 1982; Mortimore et al. 1988; Sakai et al. 1998). The lysosomal membrane can form invaginations or fingerlike protrusions to form vesicles loaded with cytoplasmic components. Once the vesicle is inside the lysosome, the membrane disappears and the sequestered material is degraded (Fig. 1). To date, the mechanisms of induction or regulation of microautophagy in mammalian cells remain unknown, and it is assumed that microautophagy participates in the continuous basal turnover of cellular components under normal cellular conditions.

*Macroautophagy* is the best-characterized form of autophagy and is frequently referred just as "autophagy." Macroautophagy is responsible for the degradation of most intracellular components as a major mechanism to maintain cellular homeostasis and appropriate balance between protein synthesis and degradation. During macroautophagy, a portion of cytosol, including entire organelles, is surrounded by an intracellular membrane to form a double-membrane organelle called *autophagosome*, which contains the sequestered components (Wang and Klionsky 2003). This vesicle fusions later with lysosomes, which provides the hydrolytic enzymes required for the total degradation of the sequestered material (Fig. 1). Under stress conditions, autophagy can be activated as part of the cellular stress-response mechanism, as it occurs, for instance, in early stages of nutrient deprivation in order to provide amino acids and other essential basic elements (He and Klionsky 2009; Ravikumar et al. 2010). In addition to this well-known role of autophagy, this pathway also participates in other cell functions such as cell differentiation, tissue remodeling, development, host-to-pathogen response, and removal of toxic or abnormal intracellular components (Mizushima et al. 2008).

The demand for autophagy activity varies among different tissues and cell types. In most tissues, macroautophagy is mainly induced upon stress conditions. Consequently, a failure in macroautophagy can result in an increased cell susceptibility to stress, since the cell is no longer able to respond properly to stress situations, but under basal conditions cell survival may not be compromised. In contrast, postmitotic cells like neurons require an effective and constitutive macroautophagy activity to maintain cell homeostasis. Consequently, any disruption of autophagy in neurons results in an accumulation of abnormal proteins within the cell and leads to neurodegeneration, even in the absence of any aggravating event such as the presence of mutant or prone-to-aggregate proteins (Hara et al. 2006; Komatsu et al. 2006; Wang and Klionsky 2003).

Macroautophagy is a complex mechanism that can be divided into different steps: induction, elongation of the limiting membrane or phagophore, cargo recognition of the substrate to be degraded, fusion of the autophagosome with lysosomes, and clearance of the cargo (Fig. 1). While macroautophagy was originally described as a nonselective step in the autophagy process, recent studies demonstrated the selectivity of the autophagic degradation process. As a result, new types of autophagy have been described according to the selective substrate involved, including *mitophagy* (i.e., selective degradation of mitochondria by autophagy) (Kim et al. 2007), pexophagy (i.e., selective degradation of peroxisomes by autophagy) (Dunn et al. 2005; Sakai et al. 1998), reticulophagy (i.e., selective degradation of endoplasmic reticulum portions by autophagy) (Bernales et al. 2007), ribophagy (i.e., selective degradation of ribosomes by autophagy) (Kraft et al. 2008), lipomacrophagy (i.e., selective degradation of lipid droplets portions by autophagy) (Singh et al. 2009), xenophagy (i.e., selective degradation of intracellular pathogens by autophagy) (Levine 2005), and aggregophagy (i.e., selective autophagic degradation of ubiquitinated aggregates).

The intraneuronal accumulation of autophagosomes and aberrant or misfolded proteins that eventually aggregate forming inclusion bodies is observed in affected brain regions of several neurodegenerative disorders and represent one of the hallmarks of these diseases, thus emphasizing the role of the autophagy system in the pathogenesis of these disorders. However, in each neurodegenerative disorders, autophagy can be affected at different step/s of the process, and therefore, the potential therapeutic approaches aimed to reestablish normal degradation activity in these diseases should differ depending on the autophagy step that is impaired.

## 3 Role of Autophagy in Neurodegenerative Disorders

Growing evidence indicates a role of autophagy failure in neurodegenerative disorders (Martinez-Vicente and Cuervo 2007; Mizushima et al. 2008; Wong and Cuervo 2010). When the intracellular degradative systems become compromised, the protein quality control of the cell fails, and as a consequence, many altered proteins (including mutant, posttranslational modified, or damaged proteins) often have the tendency to organize into complex structures like oligomers, protofibrils, and small aggregates that may be toxic to the cells. As part of the protein quality control system, cells have developed mechanisms to fight against protein misfolding, first trying to identify and repair altered proteins with chaperones (Voisine et al. 2010) and then, if necessary, eliminating the aberrant misfolded proteins through UPS or autophagy. Defects in either molecular chaperones or proteolytic systems have been identified in different neurodegenerative disorders, which have been proposed to contribute to the pathogenesis of these diseases. A failure in macroautophagy, in particular, can be dramatic for the cell since it is the only degradative mechanism able to eliminate the toxic protofibrils and aggregates once these have become insoluble. Furthermore, autophagy also contributes to the selective turnover of organelles, such as mitochondria, through the selective macroautophagy machinery. Therefore, an impairment in autophagy like the one observed in some neurodegenerative disease can lead to the accumulation of old or damaged mitochondria that become a new source of reactive oxygen species (ROS), thereby contributing to oxidative stress and aggravating the consequences of the pathologic process. Further supporting a role for autophagy in neurodegeneration, it has been reported that pharmacological or genetic inhibition of autophagy worsens the pathology and increases the number of aggregates in different cellular and animal models of neurodegeneration, while activation of macroautophagy with drugs like rapamycin can reduce the number of intracellular aggregates and ameliorates the neurodegenerative process in these experimental systems (Bove et al. 2011).

## 4 Autophagy and Parkinson's Disease

*Parkinson's disease* is a common neurodegenerative disorder of unknown origin mainly characterized by the loss of dopaminergic neurons from the substantia nigra pars compacta (SNpc) and the presence, in the affected brain

regions, of ubiquitinated intraneuronal proteinaceous cytoplasmic inclusions called Lewy bodies (LB). The presence of Lewy bodies in PD suggests that defective protein handling may contribute to the pathogenesis of the disease. The first evidence linking autophagy with PD involved the observation of accumulated autophagosomes in the SNpc of postmortem brain samples from PD patients (Anglade et al. 1997; Voisine et al. 2010; Zhu et al. 2003). Recent studies confirmed this observation (Chu et al. 2009; Dehay et al. 2010). More recently, a pathogenic autophagy impairment was observed in the MPTP mouse model of PD. MPTPtreated mice exhibit accumulations of undegraded autophagosomes and reductions in the number of functional lysosomes in SNpc dopaminergic neurons that precede cell death in this model. Lysosomal deficiency in this model was attributed to abnormal lysosomal membrane permeabilization (LMP) mediated by mitochondria-derived oxidative attack. Following LMP, the release of hydrolytic lysosomal enzymes to the cytosol can, among other consequences, abolish the lysosomal capacity of the cell, which results in an accumulation of undegraded autophagosomes that cannot be cleared by dysfunctional lysosomes (Dehay et al. 2010). Pharmacological reactivation of lysosomal-mediated degradation by rapamycin in MPTP-treated mice can restore autophagosome clearance and attenuate SNpc dopaminergic cell death in this model of PD (Dehay et al. 2010). Rapamycin also provides neuroprotection in other in vivo models of PD, such as 6-OHDA-treated mice (Santini et al. 2009), lactacystin-microinjected mice (Pan et al. 2008b), transgenic mice overexpressing wild-type (WT)  $\alpha$ -synuclein (Crews et al. 2010), or paraquat-treated drosophila (Bjedov et al. 2010). The neuroprotective effects of rapamycin in all these models were attributed, at least in part, to the activation of autophagy.

# 5 PD-Related Proteins Involved in Autophagy

Genetic studies have provided important insights into the pathological mechanism underlying PD. Among all the genes linked to familial forms of PD, many of them appear now related to autophagy pathways, thereby providing new evidences and mechanisms to potentially explain the link between autophagic failure and PD. In this section, we will discuss some of the findings linking PD-related genes/ proteins with the autophagy system.

## 5.1 $\alpha$ -Synuclein

 $\alpha$ -Synuclein is a small protein encoded by the SNCA gene. Three missense point mutations have been linked to autosomal dominant forms of PD (A30P, A53T, and E46K), as well as duplications and triplications of the gene (Chartier-Harlin et al. 2004; Ibanez et al. 2004; Kruger et al. 1998; Polymeropoulos et al. 1997; Singleton et al. 2003; Zarranz et al. 2004).  $\alpha$ -Synuclein is mostly found in neuronal presynaptic terminals, and although its exact functions remain unknown, it has

been proposed to be involved in the regulation of synaptic vesicles, neurotransmitter release, molecular chaperone activity in the SNARE complexes, and vesicle trafficking (Bonini and Giasson 2005; Burre et al. 2010; Cooper et al. 2006).  $\alpha$ -Synuclein is a major component of LB and is normally degraded by different proteolytic systems, including the UPS (Bennett et al. 1999; Imai et al. 2003), macroautophagy, and CMA (Cuervo et al. 2004; Vogiatzi et al. 2008; Webb et al. 2003; Xilouri et al. 2009). As PD progresses, alterations in UPS and autophagy systems may impair the capacity of the cell to properly handle and execute  $\alpha$ -synuclein turnover and thus contribute to the formation of protein inclusions like LB. The presence of aberrant or mutated forms of  $\alpha$ -synuclein can induce the failure of one of these proteolytic pathways and interfere with the rest of proteolytic mechanisms, given the important cross talk between all the proteolytic systems of the cell.

 $\alpha$ -Synuclein contains the CMA-targeting motif needed for the recognition by the hsc70 chaperone, which delivers the substrate to the lysosomal membrane receptor LAMP-2A in order to be translocated into the lysosomal lumen for its degradation. The two main point mutations in the  $\alpha$ -synuclein gene (A30P and A53T) that lead to familial forms of PD can affect CMA activity. When one of these α-synuclein mutants is recognized through its CMA-motif, it is targeted to lysosomes and binds LAMP-2A receptor with higher affinity than the WT protein. However, the translocation of mutant  $\alpha$ -synuclein inside the lysosome is impaired, and therefore, it cannot be degraded by CMA. In addition, mutant  $\alpha$ -synuclein remains bound to the CMA translocation complex, thereby blocking not only their own degradation but also the degradation of other CMA substrates (Cuervo et al. 2004). The pathogenic effect of mutant α-synuclein on CMA activity was also observed in neuronal cell cultures (Vogiatzi et al. 2008; Xilouri et al. 2009). Additionally, CMA failure linked to  $\alpha$ -synuclein pathogenic mutations is followed by a compensatory induction of macroautophagy, as it has been observed in other situations (Massey et al. 2006b).

Abnormal high levels of WT  $\alpha$ -synuclein are also associated with PD. In cellular models, overexpression of WT  $\alpha$ -synuclein impairs CMA activity and contributes to cell death through an effect on CMA-substrate MEF2D (Yang et al. 2009). MEF2D is a transcriptional factor required for neuronal survival that is shuttled from the neuronal nucleus to the cytoplasm, where it interacts with chaperone protein hsc70 before being targeted to lysosomes for degradation. High levels of  $\alpha$ -synuclein (like those observed in PD patients with triplications of the  $\alpha$ -synuclein gene) in  $\alpha$ -synuclein-overexpressing cellular and animal models impair CMA-mediated degradation of MEF2D, leading to the accumulation of MEF2D in the cytoplasm and subsequent neuronal death (Yang et al. 2009).

In addition to the rare  $\alpha$ -synuclein mutations and duplications/triplications of the gene (which occur in less than 2 % of patients from familial PD), the most common posttranslational modifications of  $\alpha$ -synuclein have been described in sporadic PD patients and experimental PD models (Anderson et al. 2006; McFarlane et al. 2005; Paxinou et al. 2001; Smith et al. 2005; Uversky et al. 2005; Volles and Lansbury 2003). These modifications include phosphorylation, ubiquitination, nitration, and

oxidation of  $\alpha$ -synuclein, as well as  $\alpha$ -synuclein modifications caused by its interaction with oxidized dopamine (Norris et al. 2005). Some of these posttranslational modifications of WT  $\alpha$ -synuclein can promote the formation of oligometric protofibril intermediates, which usually evolve into insoluble  $\alpha$ -synuclein fibrils, the main component of LB (Chen and Feany 2005; Conway et al. 2000; Rochet et al. 2004; Uversky et al. 2005). The ability of the modified  $\alpha$ -synuclein forms to be degraded by CMA depends on the type of modification (Martinez-Vicente et al. 2008). For instance, phosphorylation, nitration, and oxidation of  $\alpha$ -synuclein decrease the capacity of  $\alpha$ -synuclein to be translocated and eliminated through the CMA pathway, thereby increasing the amount of these  $\alpha$ -synuclein soluble forms into the cytosol and thus the possibility to interact with other molecules and become oligomers or aggregates. In contrast, dopamine-modified  $\alpha$ -synuclein behaves as the A30P and A53T mutant  $\alpha$ -synuclein, thereby remaining bound to the CMA translocation complex and blocking the degradation of other CMA substrates. The origin of this blockage seems to reside in the ability of dopaminemodified  $\alpha$ -synuclein to form oligomers on the lysosomal surface after binding as a monomer to the CMA translocation complex. The formation of these clusters on the lysosomal membrane seems to prevent the binding and translocation of other proteins. The consequences of this inability to degrade modified forms of  $\alpha$ -synuclein result in increased cytosolic concentrations of the protein, thus favoring its oligomerization. On the other hand, normal substrate proteins for CMA can no longer be turned over through this pathway and also end up accumulating inside the affected cells.

Overexpression of WT  $\alpha$ -synuclein can also inhibit macroautophagy activity, as it has been recently showed in PD models (Winslow et al. 2010). In particular, the overexpression of  $\alpha$ -synuclein causes the loss of function of Rab1, a protein involved in secretory pathways. Inhibition of Rab1a activity, in turn, causes mislocalization of Atg9, an autophagy protein essential for autophagosome formation. As a consequence, autophagosome formation is impaired, thereby resulting in an accumulation of autophagic substrates.

# 5.2 LRRK2

*LRRK2* is a large protein with several domains including, among others, a ROC domain (with GTPase activity), a COR (C-terminal of ROC) domain, a leucine-rich repeat domain (LRR), and a serine/threonine kinase domain (Gloeckner et al. 2006; Greggio et al. 2006, 2008; Greggio and Cookson 2009; Smith et al. 2006; West et al. 2005). Several mutations in this protein cause an autosomal dominant form of PD (Paisan-Ruiz et al. 2004) with pathological and clinical features very similar to late-onset sporadic PD, including the presence, in most cases, of LB. The majority of these pathological mutations are localized in the GTPase or the kinase domain, being G2019S the most common mutation present in PD cases (Dachsel et al. 2010; Paisan-Ruiz et al. 2004, 2008; Zimprich et al. 2004). Many different physiological roles for LRRK2 have been proposed, but the putative substrate/s that LRRK2

might phosphorylate in vivo remains controversial. So far, it is known that the kinase activity is required for its GTPase activity and this kinase activity depends on the formation of LRRK2 dimers. It has also been shown that LRRK2 kinase activity is required for neurodegeneration, and therefore, inhibition of its kinase activity has neuroprotective effects in PD models (Greggio et al. 2006; Smith et al. 2006).

In the last few years, LRRK2 has been suggested to play a role in numerous different cellular mechanisms. Some of the suggested functions of LRRK2 include modulation of protein translation through the direct phosphorylation of the translation initiation factor 4E-BP1 (Imai et al. 2008) and the interaction with microRNAs that regulate protein translation (Gehrke et al. 2010). In cellular models, apoptosis also seems to be regulated by LRRK2 through the direct interaction of LRRK2 with the death adaptor FADD (Ho et al. 2009). Different groups have shown that LRRK2 can also be involved in neuronal morphogenesis (MacLeod et al. 2006; Parisiadou et al. 2009; Plowey et al. 2008; Wang et al. 2008). Relevant to this chapter, other authors have implicated LRRK2 in autophagy and the lysosomal system.

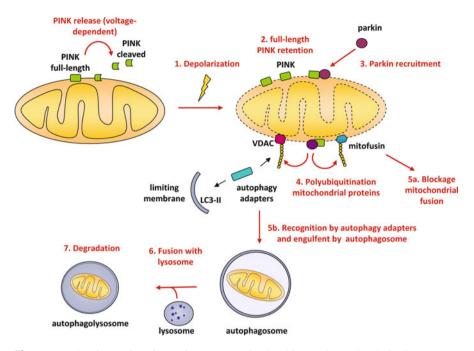
While the role of LRRK2 on autophagy is not fully understood, some authors propose that this function could be due, in part, to the role of LRRK2 on cytoskeleton dynamics and its interaction with tubulins, microtubules, and other proteins involved in trafficking and endocytosis. LRRK2 loss-of-function mutations would impair the trafficking and the cytoskeleton dynamics, thereby affecting, among other mechanisms, the lysosomal system (Dzamko et al. 2010; Gandhi et al. 2008; Gillardon 2009a, b; Sakaguchi-Nakashima et al. 2007; Shin et al. 2008). At the same time, other authors used different cellular models expressing mutant LRRK2 or LRRK2 knockdown animals to confirm the hypothesis that LRRK2 is involved in the autophagy process. These cellular and animal models exhibited an LRRK2dependent impairment of the autophagy process, thereby confirming the role of this protein in the correct functioning of autophagy (Alegre-Abarrategui et al. 2009; Ferree et al. 2011; Gomez-Suaga et al. 2012; Ramonet et al. 2011; Tong et al. 2010, 2012). More recently, a novel PD cellular model based on the generation of dopaminergic neurons derived from induced pluripotent stem cells (iPSC) obtained from fibroblasts of patients carrying the G2019S LRRK2 mutation exhibited the typical autophagic alterations observed in postmortem brain samples from PD patients, such as an accumulation of autophagic vacuoles and defects in autophagosome clearance (Sanchez-Danes et al. 2012). Taken together, these data confirm the implication of LRRK2 in the autophagy process. However, the mechanisms underlying this effect and the consequences of PD-linked mutations of LRRK2 on this function remain unknown.

## 5.3 Parkin and PINK1

Autophagic failure in PD has recently been confirmed through the newly identified roles of parkin and *PINK1* proteins in maintaining mitochondria homeostasis by mediating mitochondrial trafficking and autophagic clearance of dysfunctional

mitochondria. *Parkin* is a component of a multiprotein E3 ubiquitin ligase and PINK1 is a serine/threonine protein kinase that localizes in the mitochondria and is involved in stress-induced mitochondrial dysfunction. Recently, both proteins have been proposed to belong to the mitochondrial quality control system, working together to promote degradation of dysfunctional mitochondria by mitophagy (Dagda et al. 2009; Narendra et al. 2008; Vives-Bauza et al. 2009; Whitworth and Pallanck 2009).

Loss-of-function mutations in PINK1 or parkin are associated to autosomal recessive familial PD. These mutations result in mitochondrial dysfunction with impaired mitochondrial turnover that leads to cell death. Indeed, it has recently been shown that PD-linked mutations in parkin can cause defects in the mitophagy process (Geisler et al. 2010; Lee et al. 2010b; Matsuda et al. 2010; Narendra et al. 2010b; Vives-Bauza et al. 2010). In addition, PD-linked mutations in PINK1 cause defective parkin recruitment to mitochondria and, consequently, mitophagy efficiency is affected (Geisler et al. 2010; Narendra et al. 2010b; Ziviani and Whitworth 2010; Vives-Bauza et al. 2010). PINK1is localized in mitochondria where it is cleaved to a shorter form that is no longer retained in mitochondria and is released to the cytosol, resulting in low mitochondrial levels of full-length PINK. This PINK1 cleavage is membrane-voltage dependent, and after depolarization of the mitochondrial membrane, the cleavage is inhibited, resulting in an accumulation of full-length PINK1in the mitochondria (Narendra et al. 2010b) (Fig. 2). The retention of PINK1 in damaged mitochondria recruits parkin to mitochondria which, through its E3 ubiquitin ligase activity, can polyubiquitinate different membrane proteins like VDAC1 and mitofusin (Geisler et al. 2010; Narendra et al., 2010b; Ziviani et al. 2010; Gegg et al. 2010). Polyubiquitination of mitofusin inhibits mitochondrial fusion (Tanaka et al. 2010; Ziviani and Whitworth 2010), thereby preventing the fusion of damaged mitochondria with healthy ones, while polyubiquitination of VDAC1 is required for mitochondrial clearance by mitophagy (Fig. 2). After VDAC1 polyubiquitination by parkin, depolarized mitochondria is tagged and autophagic adapters can bind the mitochondria to be sent to the autophagosome, where it will be degraded after fusion with lysosomes. A potential role for the autophagic adapter p62 in this parkinmediated mitophagy process is still controversial. p62 can bind both ubiquitin proteins, by its ubiquitin-associated (UBA) domain, and also LC3, through its LC3-interacting region (LIR) domain, thus acting as a recruiter of ubiquitin-tagged substrates that are sent to autophagic limiting membrane, where LC3 is one of the main components (Pankiv et al. 2007). The autophagosome is then formed and the substrate is eliminated by macroautophagy. Some authors have reported that this PINK/parkin-mediated process is p62-dependent and that the damaged mitochondria tagged by ubiquitination requires the binding of p62 to be eliminated (Ding et al. 2010; Geisler et al. 2010; Okatsu et al. 2010). However, other studies have found that mitophagy is not p62-dependent and probably other autophagic adapters can display this function (Lee et al. 2010a; Narendra et al. 2010a; Okatsu et al. 2010).



**Fig. 2** *PINK/parkin-mediated mitophagy.* Upon mitochondria membrane depolarization, PINK1 full-length protein is retained in mitochondria membrane and recruits parkin protein. Once parkin is localized in the mitochondria membrane, they can polyubiquitinate different membrane proteins like VDAC1 and mitofusin. Polyubiquitinization of mitofusin inhibits future mitochondrial fusion with other healthy mitochondria, while polyubiquitinization of VDAC allows the interaction with autophagic adapters that will trigger the engulfment of the depolarized mitochondria by the autophagosome and its subsequent degradation inside the lysosomes

## 5.4 DJ-1

Mutations in the *DJ*-1 gene result in a rare cause of autosomal recessive PD (Bonifati et al. 2003; Wong and Cuervo 2010). The physiological role of DJ-1 is still not fully understood, but it has been implicated in different biological functions like maintaining mitochondria integrity and acts as a redox-sensitive chaperone with antioxidant properties (Blackinton et al. 2009; Canet-Aviles et al. 2004; Irrcher et al. 2010; Junn et al. 2009; Kim et al. 2005; Meulener et al. 2005; Taira et al. 2004; Wong and Cuervo 2010). Most of these studies were performed in experimental in vivo and in vitro models of DJ-1 knockdown and showed that loss of DJ-1 function is linked to mitochondria depolarization and fragmentation, hypersensitivity to oxidative stress, and reduction of basal autophagy (Krebiehl et al. 2010). Recently, DJ-1 has also been linked to mitophagy by working together with parkin and PINK1 in the elimination of damaged mitochondria after oxidative stress (Thomas et al. 2011). The mechanism of action of DJ-1 on mitophagy remains unclear, but deficiency in DJ-1 leads to a similar mitochondrial phenotype as the one observed in

PINK1- and parkin-deficiency models (Thomas et al. 2011). In addition, DJ-1 overexpression is able to protect against oxidative stress in PINK1-deficient cells (Thomas et al. 2011). Overexpression of parkin or PINK1 can rescue the mito-chondrial alterations occurring in DJ-1-knockdown cells, thus suggesting a potential role for DJ-1 in PINK1/parkin-mediated mitophagy.

## 5.5 UCH-L1

Another potential link between PD pathogenesis and autophagy is the role of UCH-L1 protein in the CMA pathway. UCH-L1 (ubiquitin C-terminal hydrolase L1) is expressed in neurons and is associated with familial PD forms. This protein physically interacts with CMA receptor LAMP-2A at the lysosomal membrane. the UCH-L1 protein carries the PD-linked mutation I93M When or a posttranslational carbonyl modification reported in sporadic forms of PD, the interaction of the protein with the CMA translocation complex is increased, thereby preventing the CMA-mediated degradation of other substrates, as previously observed with some forms of  $\alpha$ -synuclein (Kabuta et al. 2008). Interestingly, the UCH-L1 polymorphism S18Y may have opposite effects, being protective against the development of sporadic PD (Satoh and Kuroda 2001; Elbaz et al. 2003; Carmine et al. 2007). Experimental in vitro and in vivo PD models experimentally confirmed a neuroprotective effect of the S18Y polymorphism (Kyratzi et al. 2008; Xilouri et al. 2012). While the basis for this neuroprotective effect is still controversial, it may involve the antioxidant function of this UCH-L1 variant.

## 5.6 ATP13A2

The ATP13A2/PARK9 gene encodes the protein ATP13A2, a lysosomal type 5 P-type ATPase present in lysosomal membranes, that is linked to autosomal recessive, levodopa responsive, nigrostriatal-pallidal-pyramidal neurodegeneration (Kufor-Rakeb Syndrome) as well as to some juvenile and young-onset forms of PD parkinsonism (Di Fonzo et al. 2007). The exact mechanism underlying the cellular functions of human ATP13A2 remains unknown, as well as its role in PD. Recently, it was shown that mutations/loss of ATP13A2 in human fibroblasts from patients with Kufor-Rakeb syndrome or in mouse primary neurons and dopaminergic cell lines leads to impaired lysosomal degradation capacity (Dehay et al. 2012; Usenovic et al. 2012a). ATP13A2 protein plays an important role in lysosomal membrane integrity, and mutations present in the ATP13A2/ PARK9 gene cause instability of the lysosomal membrane, impaired lysosomal acidification, decreased proteolytic capacity, and diminished lysosomal-mediated clearance of autophagosomes (Dehay et al. 2012). This lysosomal dysfunction results in accumulation of  $\alpha$ -synuclein and toxicity in primary cortical neurons (Usenovic et al. 2012a). In addition, silencing of endogenous  $\alpha$ -synuclein attenuated the toxicity in ATP13A2-depleted neurons, suggesting that loss of ATP13A2

mediates neurotoxicity at least in part via the accumulation of  $\alpha$ -synuclein (Usenovic et al. 2012a). Identification of ATP13A2-interacting proteins by yeast two-hybrid system provide insights into ATP13A2 function and suggest a role in vesicular trafficking and fusion (Usenovic et al. 2012b). Loss of function of this protein could mediate  $\alpha$ -synuclein aggregation and toxicity and explain the link between ATP13A2 mutations observed in patients with Kufor-Rakeb syndrome and neurodegeneration. In addition, loss of ATP13A2 function in cellular models was reported to disrupt autophagy activity and affect, in particular, mitochondrial quality control, thereby leading to impairment of mitochondrial turnover and increase of ROS production (Gusdon et al. 2012).

## 5.7 Glucocerebrosidase (GBA)

Loss-of-function mutations in the *GBA* gene are associated to *Gaucher's disease* (GD) and are also linked to an increased risk of PD and other related  $\alpha$ -synucleinopathies (Neumann et al. 2009). *Glucocerebrosidase* protein (GCase) is a lysosomal enzyme responsible for the breakdown of the glycolipid glucocerebrosidase into glucose and ceramide inside lysosomes. GD is an autosomal recessive lysosomal storage disease caused by loss-of-function mutations of the GCase enzyme that results in accumulation of GCase substrates inside the lysosomes, especially in macrophages. GD is clinically classified in three groups: type 1 or non-neuropathic, type 2 or acute neuropathic, and type 3 or chronic neuropathic. Interestingly, patients with GD type 1, the non-neuropathic variant, have increased propensity to develop PD or related LB diseases (DePaolo et al. 2009). In turn, patients with PD have increased frequency of GBA mutations (DePaolo et al. 2009).

Although the mechanism underlying the association of GBA with PD remains unknown, several groups have proposed that impaired autophagy, aggregate formation, and  $\alpha$ -synuclein metabolism may link GBA mutations with the pathogenesis of PD (Westbroek et al. 2011). In particular, some studies propose that alterations in the lysosomal system secondary to the presence of mutant GCase might connect PD and GD (Sun et al. 2010; Vitner et al. 2010). In cell culture models, the GBA mutants with gain-of-toxic function promote  $\alpha$ -synuclein accumulation, an effect that can be reverted by autophagy induction with rapamycin (Cullen et al. 2011). Furthermore, patients with GD and PD exhibit LB positive for GCase, which suggests that  $\alpha$ -synuclein oligomers may sequester mutant misfolded GCase within LB during the progression of the disease (Cullen et al. 2011). These results indicate that loss of GBA creates a positive feedback loop of reduced lysosomal function and  $\alpha$ -synuclein accumulation that ultimately leads to neurodegeneration (Dawson and Dawson 2011; Mazzulli et al. 2011; Sardi et al. 2011).

Additional work is necessary to further identify the mechanisms underlying the connection between PD and GD, as other not-yet-identified factors or proteins may probably play a role since only a small percentage of GD patients and carriers with GBA mutations actually develop PD.

## 6 Conclusion

Mounting evidence indicate a role for autophagy dysfunction in neurodegenerative diseases, including PD. The potential involvement of autophagy alterations in PD has been reinforced in recent years by the finding that many of the mutated nuclear genes associated with familial forms of PD either directly or indirectly affect the normal activity of autophagy pathways. These findings might help to explain the pathogenesis of these diseases and most importantly lay the ground for the development of new therapeutic strategies based on the modulation of autophagy pathways for this group of devastating, currently incurable, diseases.

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# Dopaminergic Neurons in Parkinson's Disease

Rui Daniel Prediger, Mariza Bortolanza, Ana Carolina de Castro Issy, Bruno Lopes dos Santos, Elaine Del Bel, and Rita Raisman-Vozari

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R.D. Prediger

M. Bortolanza • A.C. de Castro Issy • E. Del Bel

Departamento MFP-Fisiologia, Faculdade de Odontologia de Ribeirão Preto, Universidade de São Paulo (USP), Center for Interdisciplinary Research on Applied Neurosciences (NAPNA), Ribeirão Preto, SP, Brazil

e-mail: marizabortolanza@yahoo.com.br; carol\_issy@hotmail.com; eadelbel@forp.usp.br

B.L. dos Santos

R. Raisman-Vozari (🖂)

Laboratório Experimental de Doenças Neurodegenerativas, Departamento de Farmacologia, Universidade Federal de Santa Catarina (UFSC), Florianópolis, SC, Brazil e-mail: ruidsp@hotmail.com

Department of Neurosciences and Behavior Sciences, Medical School of Ribeirão Preto, University of São Paulo, Ribeirão Preto, SP, Brazil e-mail: bruls4@usp.br

INSERM UMR 975 (ex U679), CNRS UMR 7225, Université Pierre et Marie Curie (UPMC), Centre de Recherche de l'Institut du Cerveau et de la Moelle épinière (CRICM), Thérapeutique Expérimentale de la neurodégénérescence, Hôpital de la Salpêtrière, Paris, France e-mail: ritaraisman@gmail.com

#### Abstract

The discovery of dopamine as a brain neurotransmitter by Arvid Carlsson and colleagues about 50 years ago contributed to better understand some of the brain diseases. Some of the drugs that are most widely used in neurologic and psychiatry illnesses, such as levodopa and antipsychotic drugs, act on dopaminergic mechanism. The discovery that the motor impairments of Parkinson's disease patients improved after restoring the physiological levels of striatal dopamine with levodopa attracted the attention of the neuroscience community for the role of this neurotransmitter in motor and brain functions. In the last decades, the knowledge has also been challenged by evidence that Parkinson's disease also affects cognitive and affective functions. Shortly after the introduction of levodopa as a therapy, a complex set of secondary phenomena such as dyskinesia was observed following repeated administration of the dopamine precursor. Information of dopaminergic cells and circuits has been enriched by findings obtained with several and highly sensitive technology in cellular biology, with sophisticated behavioral analyses of transgenic animals and functional neuroimaging. The present chapter attempts to review results reported in different clinical studies and animal models to provide a comprehensive picture of the role of dopamine in Parkinson's disease. Treatments have successfully been translated from preclinical to pharmacotherapeutic arsenal increasing clinical settings.

Keywords

Basal ganglia • Dopamine • Dopamine receptors • Dopaminergic drugs • Levodopa • Mesocortical system • Mesolimbic system • Nigrostriatal system • Nitric oxide • Parkinson's disease

List of Abbreviations				
3-OMD	3-O-methyldopa			
6-OHDA	6-hydroxydopamine			
AIMs	Abnormal involuntary movements			
AS	Alpha-synuclein			
CNS	Central nervous system			
COMT	Catechol-O-methyltransferase			
DARPP-32	Dopamine- and cAMP-regulated phosphoprotein, Mr 32 kDa			
DAT	Dopamine transporter			
DLB	Dementia with Lewy body			
DMV	Dorsal motor nucleus of vagus			
DYN	Dynorphin			
ENK	Enkephalin			
FS	Fast spiking			
GPCRs	G protein-coupled receptors			
GPe	External segment of the globus pallidus			
GPi	Internal segment of the globus pallidus			

ית די	
iLBP	Incident Lewy body pathology
IrtZ	Intermediate reticular zone
i.v.	Intravenous
LB	Lewy bodies
LC	Locus coeruleus
L-DOPA	L-3,4-dihydroxyphenylalanine
LN	Lewy neurites
LTS	Low-threshold spiking
MAO-B	Monoamine oxidase-B
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine
MSN	Medium spiny neuron
NPY	Neuropeptide Y
PD	Parkinson's disease
PET	Positive emission tomography
PV	Parvalbumin
SN	Substantia nigra
SNc	Substantia nigra pars compacta
SNr	Substantia nigra pars reticulata
SOM	Somatostatin
SP	Substance P
SPECT	Single-photon emission computed tomography
STN	Subthalamic nucleus
TH	Tyrosine hydroxylase
VMAT2	Vesicular monoamine transporter 2
VTA	Ventral tegmental area
	U U

## 1 Introduction

At the beginning of the nineteenth century, when James Parkinson published the book entitled "*Essay on the Shaking Palsy*" (Parkinson 2002), life expectation was no longer than 45 years old. The "shaking palsy," later named Parkinson's disease (PD) by Charcot, affects mainly aged people. Nowadays, the prevalence of PD in industrialized countries is generally estimated at 0.3 % of the entire population and about 1 % in people over 60 years of age (Nussbaum and Ellis 2003). Age disease incidence ranges from 110 to 330 per 100,000 individuals over 50 years old, and its incidence increases with increasing age throughout the life span (Mayeux 2003). Now, in the beginning of the twenty-first century, life expectancy is near 80 years old, and the incidence rate of PD increases to 400–500 individuals per 100,000 annually (Mayeux 2003). Therefore, there are much more parkinsonian patients nowadays than two centuries ago, and with the increasing age of the general population, the prevalence of PD will rise constantly in the future.

PD is characterized clinically by bradykinesia, rest tremor, muscle rigidity, and postural reflex impairment. The discovery that the motor impairments of PD patients improved after restoring the physiological levels of striatal dopamine with levodopa attracted the attention of the neuroscience community for the role of this neurotransmitter in motor functions (Olanow and Tatton 1999). Since dopamine is the main modulator of the basal ganglia, their components have long been seen as part of the motor system (Graybiel et al. 1994). In the last decades, this view has been challenged by evidence that PD also affects cognitive and affective functions (Dubois and Pillon 1997; Chaudhuri et al. 2006; Da Cunha et al. 2009). In addition, dysfunction of dopamine neurotransmission has been implicated in many other psychiatric and neurological diseases, such as attention deficit hyperactivity disorder (ADHD), drug addiction, Tourette syndrome, schizophrenia, and obsessive-compulsive syndrome. Furthermore, a growing body of evidence has been accumulated supporting the view that some kinds of cognitive processes, such as instrumental learning, decision-making, and other executive functions, depend critically on the dopamine signaling (see Da Cunha et al. 2009).

The present chapter attempts to review results reported in different clinical studies and animal models to provide a comprehensive picture of the role of dopamine signaling in PD.

# 2 Dopamine Neurotransmission System: Pathways, Receptors, and Signaling

A detailed review of the history and neurobiology of the dopamine signaling is beyond the scope of this chapter and can be found in very interesting reviews by Hornykiewicz (2002), Iversen and Iversen (2007), and Fahn (2008). The neurotransmitter dopamine plays a crucial role in motor, motivational, and rewardrelated functions of the central nervous system (CNS). Our understanding of neuropsychiatric illnesses and the modern psychopharmacology was greatly influenced by the discovery of dopamine as a neurotransmitter in the brain by Arvid Carlsson and colleagues about 50 years ago. Some of the drugs that are most widely used in the treatment of neurologic and psychiatry illnesses, such as levodopa, methylphenidate, and antipsychotic drugs, act on dopaminergic mechanisms. Therefore, dopamine neurotransmitter system has been one of the major and fertile fields of behavioral neuroscience and its research culminated in the award of the Nobel Prize in 2000 to three scientists, Carlsson, Greengard, and Kandel, for their important contributions to this field.

Dopamine was the last catecholamine discovered, and Sir Henry Dale, in 1952, suggested dopamine as the current name of 3,4-dihydroxyphenethylamine, also known as 3-hydroxytyramine. Its biosynthesis begins with dietary amino acid tyrosine actively transported into the brain. Tyrosine hydroxylase (TH), the rate-limiting enzyme in catecholamine synthesis, is regulated by multiple mechanisms

(Molinoff and Axelrod 1971). Rapid activation of TH activity occurs via its phosphorylation by several protein kinases, including protein kinase A,  $Ca^{2+}$ / calmodulin-dependent protein kinase II, and protein kinase C. Possibly, its phosphorylation induces conformational change that results in a lower affinity of TH for catecholamines that trigger end-product inhibition of this enzyme. The end result is an increase of TH catalytic activity (Nestler et al. 2009). Several drugs can up- or downregulate TH expression, and extracellular stimuli may induce longer-term changes in TH activity through transcriptional regulation of its gene.

Dopamine is synthesized in the cytosol and transported into synaptic vesicles via the vesicular monoamine transporter 2 (VMAT2). From the synaptic vesicles, dopamine is released in a quantal manner when a nerve impulse traveling down the axon reaches the nerve terminal. Once in the synaptic cleft, dopamine can act on both postsynaptic and presynaptic dopamine receptors. Dopamine could be metabolized to homovanillic acid (HVA); 3,4-dihydroxyphenylacetic acid (DOPAC) via monoamine oxidase (MAO), an enzyme considered to be mostly intraneuronal; and 3-methoxytyramine (3-MT) via catechol-*O*-methyltransferase (COMT), an enzyme considered mostly extraneuronal. Entacapone and tolcapone are reversible COMT inhibitors which have been approved for clinical use in patients with PD as adjuvant to levodopa, significantly improving the clinical benefits of this therapy (Bonifácio et al. 2007).

Dopamine signal ends by spontaneous amine diffusion, local enzymatic degradation, and amine reuptake in the presynaptic neurons by dopamine transporter (DAT) (Cragg and Rice 2004; Nestler et al. 2009; Eriksen et al. 2010). This dopamine transporter seemingly play a key role in the dopamine transmission if we consider the behavioral consequences of their inhibition with blocking agents known to be powerful psychostimulants or drugs of abuse, as amphetamine or cocaine (see Nestler et al. 2009; Fig. 1).

Due to their diverse anatomy and disparate functions, different populations of dopamine neurons have been specifically implicated in a variety of neurological disorders. Biochemical characterization of dopamine in the brain revealed two families of dopamine receptors: D1-like receptors that also comprises D5 receptor and are able to stimulate adenylyl cyclase and increase cAMP production and D2-like receptors that comprises D3 and D4 receptors and are negatively coupled to adenylyl cyclase activity and inhibit cAMP production. In mammals, all of these receptors belong to the superfamily of GPCRs with seven transmembrane domains. D1 family is coupled to stimulatory  $G_s$  or the related  $G_{olf}$ , and the D2 family is coupled to inhibitory  $G_i/G_o$  (for review, see Herve and Girault 2005). D1 receptor activation stimulates the cAMP increase leading to the activation of protein kinase that have a broad array of cellular targets, including transcription factors, voltage-dependent ion channels, and glutamate receptors (Svenningsson et al. 2004).

Dopaminergic neurons and pathways were first described using classical histofluorescence techniques during the 1960s and 1970s (Carlsson et al. 1962). Knowledge of dopaminergic cells and circuits has been enriched by findings

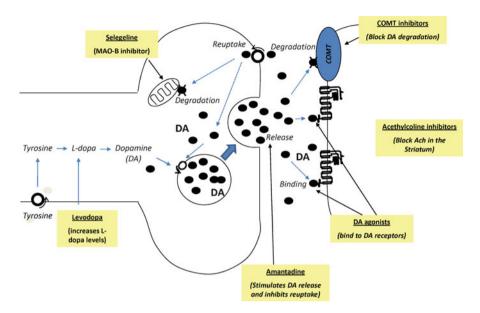
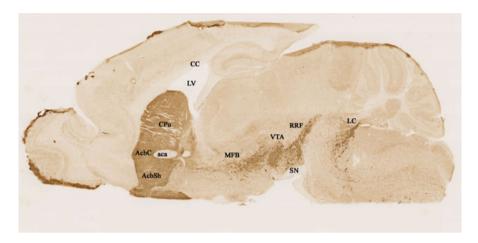


Fig. 1 Dopamine synthesis, release, signaling, and reuptake and pharmacological approaches currently available for Parkinson's disease. Dopamine synthesis originates from the amino acid tyrosine, which is converted to DOPA by the enzyme tyrosine hydroxylase (TH) (a rate-limiting enzyme). Subsequently, DOPA is decarboxylated, by the enzyme L-aromatic amino acid decarboxvlase (DOPA decarboxylase), to form dopamine. L-dopa that is used in the Parkinson's disease treatment bypasses the rate-limiting enzymatic step and is decarboxylated by DOPA decarboxylase to form dopamine. The most significant mechanism by which the synaptic actions of dopamine is terminated is the reuptake into the nerve terminal via specific dopamine transporter (DAT) expressed on the membrane of presynaptic terminal. Dopamine is also catabolized by the enzymes monoamine oxidase (MAO) and catechol-O-methyltransferase (COMT). Possibly, MAO exists in mitochondria within dopaminergic nerve terminals, on extracellular space, and also in glia proximate to dopamine synapses. After its synthesis, dopamine acts on pre- and postsynaptic receptors from D1 or D2 dopamine receptor families, which, respectively, activates Gs protein and increases cAMP production or Gi protein and decreases cAMP production. Generally drugs including L-DOPA, carbidopa, amantadine, and COMT and MAO-B inhibitors revert temporarily the dopamine dysfunction caused by dopaminergic neurodegeneration in the basal ganglia of Parkinson's disease patients

obtained with several and highly-sensitive techniques, as immunohistochemistry (mainly to dopamine essential synthetic enzyme TH), cellular biology, transgenic mice or conditional mutants, functional anatomy, electrophysiological studies, sophisticated behavioral analyses, and functional neuroimaging. Altogether, these studies have also led to new concepts of healthy and pathological functions of dopaminergic circuits at the molecular and more integrated levels (Björklund and Dunnett 2007).

In the ventral mesencephalic tegmentum, dopaminergic cell bodies are very well organized in the following groups: the substantia nigra pars compacta (SNc) (A9),



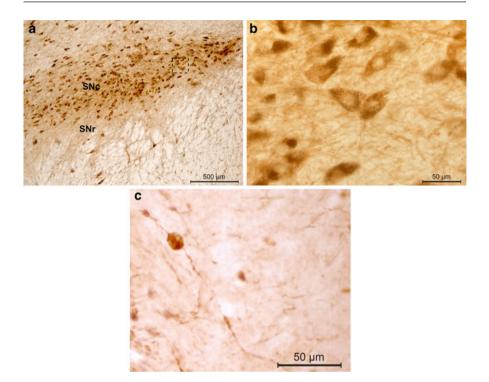
**Fig. 2** Rat brain sagittal view (adult, *Wistar*) of the tyrosine hydroxylase immunocytochemistry showing the distribution of dopaminergic neurons. Abbreviations: *aca* anterior comm, anterior part, *AcbC* accumbens nucleus, core, *AcbSh* accumbens nucleus, shell, *cc* corpus callosum, *CPu* caudate putamen (striatum), *LC* locus coeruleus, *LV* lateral ventricle, *MFB* medial forebrain bundle, *RRF* retrorubral field, *SN* substantia nigra, *VTA* ventral tegmental area

the ventral tegmental area (VTA-A10), and the retrorubral area (A8) (Fig. 2). Dahlström and Fuxe (1964) defined the nomenclature of dopaminergic system in the rat midbrain: A8, A9 (substantia nigra), and A10 cell groups. The classification is nonetheless widely used.

Ascending dopaminergic pathways in the mammalian CNS can be simplified divided into three major systems. The two largest groups are located in the midbrain: the nigrostriatal system that originates in the substantia nigra (A9) and innervates predominantly the dorsal striatum and the mesocortical and mesolimbic systems that arise from the VTA and project to the cortical or several limbic areas (Bjorklund and Lindvall 1984; Krimer et al. 1997). The cell group of retrorubral area that forms a dorsal and caudal extension of the VTA contains cells that project to striatal, limbic, and cortical areas (Bentivoglio and Morelli 2005).

## 2.1 Nigrostriatal System

Since the first detailed studies of the SN by Cajal (1911), two main subdivisions were recognized. The SNc is characterized by densely packed neurons, and the substantia nigra *pars reticulata* (SNr) is marked by sparse dopaminergic cells, enmeshed in dopaminergic dendrites (Fig. 3). Most of the properties of dopamine neurotransmission in vivo are known from studies of the nigrostriatal dopamine pathway extending from the SNc to the dorsal striatum through the median



**Fig. 3** Tyrosine hydroxylase immunocytochemistry of the regions of the human brain presenting positive reaction in the substantia nigra *compacta* (**a**, **b**; SNc), *reticulata* (**a** SNr,), and the striatum (**c**; caudate). Observe a high density of TH-positive neurons in the SNc, in contrast to sparse distribution in the SNr. TH- positive neurons are detected in the human and non human -primate striatum and and also in mice

forebrain bundle. Somato-dendritic release of dopamine in the SNc and axonal dopamine release in the dorsal striatum are both necessary for the expression of basal ganglia-mediated motor behaviors and various cognitive functions. In both rat and monkey, the striatal dopamine innervation is derived not only from SNc but also from cells located in the lateral VTA and retrorubral area. For this reason, the term "mesostriatal dopamine pathway" sometimes is used to include all components of the midbrain dopamine system projecting to the striatum (Björklund and Dunnett 2007).

The identification of cells of the SN as dopaminergic and of the dopaminergic innervation of the striatum through the nigrostriatal tract is inseparably intertwined with methodological achievements in experimental and chemical neuroanatomy in the 1960s and 1970s. These discoveries were intrinsically correlated to the histopathology of the midbrain dopaminergic system in PD in the 1960s (Bentivoglio and Morelli 2005).

## 2.2 Mesocortical and Mesolimbic Systems

Both of these systems connect the ventral tegmental dopamine cells (A8, A10) to prefrontal and cingulate cortex or to nucleus accumbens, amygdaloid complex, hippocampus, and piriform cortex. Ventral tegmental dopaminergic neurons play a critical role in motivation, reward-related behavior, attention, and multiple forms of memory (Nestler et al. 2009).

A third population of midbrain dopamine neurons resides in the retrorubral field and projects primarily to the dorsal striatum and the pontomedullary reticular formation; it is thought to play a role in orofacial movements. There are also connections between the retrorubral field and SN/VTA dopamine neurons. There are at least four groups of dopamine neurons in the hypothalamus that are intimately involved in neuroendocrine, hormonal, and arousal processes. Finally, there are populations of dopaminergic amacrine cells in the retina that contribute to neural adaptation to light and also some dopamine cells in the olfactory bulb (Nestler et al. 2009).

The striatum receives input from a number of regions, including the motor, sensory, and frontal cortex, integrates this information under the modulatory influence of dopamine, and uses this information to guide motor behavior. A cardinal feature of the striatum – the main integrator of cortical and thalamic information reaching the basal ganglia – is its dense dopamine innervation, which arises from the midbrain nigrostriatal pathway. Nigrostriatal dopaminergic neurons form highly branched networks within their striatal targets. The full extent of this arborization was elegantly demonstrated by Matsuda and colleagues (2009). A single dopaminergic neuron innervated up to 5.7 % of the neostriatum, while adjacent input dopaminergic neurons showed high degrees of overlapping with targets. It was calculated that a single postsynaptic medium spiny neuron (MSN) could be influenced by up to 194 dopaminergic inputs. As a result of this redundancy, clinical signs of PD only develop after up to 70 % of SNc neurons are lost. Each striatal MSN receives input from several thousand different cortical neurons on its spines. The input from each synapse is weak and many inputs are needed to discharge the neuron (Rothwell 2011).

Anatomically, striatal cells fall into two main classes: (a) spiny projection neurons and (b) aspiny interneurons. Spiny projection neurons, also known as medium spiny neurons (MSNs), represent the vast majority of striatal neurons. MSNs receive glutamatergic inputs from cortex and thalamus that terminate predominantly on spines (Kemp and Powell 1971). In addition, they are a main target of midbrain dopaminergic neuron axons that form synapses on MSN dendrites and spine necks (Smith et al. 1994).

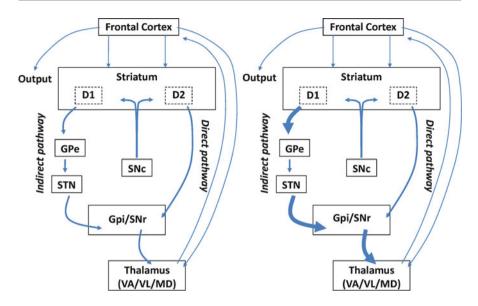
One of the major targets of this innervation is the principal neuron of the striatum: GABAergic spiny projection neurons. Spiny projection neurons constitute as much as 90 % of the striatal neuron population. They can be divided into two approximately equally sized populations based on axonal projections. One population – **direct pathway spiny projection neurons** – projects axons to the nuclei at the interface between the basal ganglia and the rest of the brain,

whereas the other population, **indirect pathway spiny projection neurons**, projects only indirectly to the interface nuclei. The dopamine D1 receptor is expressed selectively by direct pathway spiny projection neurons. Also, **direct pathway** spiny projection neurons express high levels of substance P (SP) and dynorphin (DYN). The dopamine D2 receptor is expressed by **indirect pathway** spiny projection neurons, which also express enkephalin (ENK). These two G protein-coupled receptors (GPCRs) have distinctive intracellular signaling cascades and targets, leading to fundamentally different cellular responses to extracellular dopamine (see below).

There are four well-characterized classes of striatal interneuron: cholinergic interneuron, parvalbumin-expressing GABAergic interneuron, calretinin-expressing GABAergic interneuron, and neuropeptide Y (NPY)/nitric oxide-expressing GABAergic interneuron (Tepper et al. 2008). Together, these interneurons constitute 5–10% of all striatal neurons. One of them, the cholinergic interneuron, co expresses dopamine D2 and D5 receptors (Bergson et al. 1995) and modulates both spiny projection neurons populations through muscarinic receptors (Yan and Surmeier 1996). Two other prominent interneurons, the somatostatin (SOM)/NPY-expressing GABAergic interneuron and the parvalbumin (PV) GABAergic interneuron, express dopamine D5 receptors (Centonze et al. 2003). The PV interneuron is strongly innervated by globus pallidus neurons that express dopamine D2 receptors (Bevan et al. 1998), creating a microcircuit that is influenced by both dopamine D1 and D2 receptors.

According to the classic model of basal ganglia function (see Fig. 4; DeLong 1990), movements occur during pauses in the tonic inhibitory activity of the basal ganglia output interface, generated by activity in the direct pathway. This model had its origins in neurophysiologic studies showing that corticostriatal activation of the direct striatonigral pathway results in pauses of the tonic activity of GABAergic neurons in the SNr (Chevalier et al. 1985). The opposing role of the indirect pathway in suppressing movements was originally proposed on the basis of studies in animal models of PD. Mink (2003) reviewed the classic model, proposing that the activity of direct and indirect pathways is coordinated to select particular motor programs and to inhibit competing motor programs. This model predicts that during ongoing behavior, there will be increased activity in neuronal ensembles that are part of direct and indirect pathway circuits, rather than one or the other. The execution of a movement sequence would then generate a complex pattern of activity in specific neuronal ensembles. Recent work using optogenetic or genetic approaches supports the general tenets of the model (Hikida et al. 2010; Kravitz et al. 2010). Although this model has proven to be of considerable clinical value, it fails to account for the great diversity and complexity of the decision-making process in action selection (Cisek and Kalaska 2010).

Dopamine acted by stimulating the MSNs of the direct and inhibiting those of the indirect pathway (Albin et al. 1989; Gerfen et al. 1990; Gerfen and Surmeier 2011). These opposing effects were suggested to depend on the expression of dopamine D1 receptors in the MSNs of the direct pathway and of dopamine D2 receptors in the MSNs of the indirect pathway. These receptors have distinctive



**Fig. 4** Overview of schematic representation of the main basal ganglia-thalamocortical circuits (left). Normal status: the two output routes ("indirect" and "direct") are in balance at the level of the output structures (the GPi and the SNr) (right). Neurodegenerative changes in the central nervous system in Parkinson's disease (PD): depletion of dopamine in the striatum leads to imbalance in the two output routes and suppression of thalamocortical activity. The thickness of the arrows indicates the level of activity in the pathways. For some time, the symptoms of PD have been described as a loss of balance between the direct and indirect striatal output pathways. The indirect pathway is attenuated primarily by D2 receptor stimulation, whereas the direct pathway is activated via D1 receptor stimulation. With normal tonic/phasic balance, cortical activation of the indirect pathway (attenuated by tonic D2 stimulation) is in balance with other presumably motor cortical inputs influencing the direct pathway (augmented by phasic D1 stimulation). However, in PD, this balance is disrupted, with the loss of DA resulting in hyperactivity (with akinesia) in the indirect pathway and depression (with loss of voluntary movement) within the direct pathway. The model predicts that neuronal firing in the STN and GPi is increased in the parkinsonian state, leading to excessive inhibition of brain stem and thalamocortical neurons with the development of parkinsonian motor features. In contrast, the model proposes that dyskinesia is related to the decreased firing in the STN and GPi, with reduced inhibition of thalamic and cortical motor regions. Abbreviations: D dopamine, GPe external segment of globus pallidus, GPi internal segment of globus pallidus, MD mediodorsal nucleus, SNr reticular part of the substantia nigra, SNc pars compacta of the substantia nigra, STN subthalamic nucleus, VA/VL ventral anterior/ ventral lateral thalamic nuclei (Modified from Hauser 2009)

intracellular signaling cascades and targets, leading to fundamentally different cellular responses to extracellular dopamine (Gerfen and Surmeier 2011). Also, direct pathway expresses high levels of substance P and dynorphin, whereas indirect pathway expresses enkephalin. Direct and indirect pathways are representing on Fig. 4.

Dopamine stimulation of D1 receptors acts via cAMP and protein kinase A to phosphorylate dopamine- and cAMP-regulated phosphoprotein, Mr 32 kDa

(DARPP-32), an important mediator of cAMP signaling and striatal excitability. The DARPP-32 signaling pathway has a central role in mediating signal transduction within MSNs in the striatum, and the function of this phosphoprotein depends on its relative state of phosphorylation at two main regulatory sites, threonine 34 (T34) and threonine T75 (T75). When DARPP-32 is phosphorylated at T34 by protein kinase A, it becomes a potent inhibitor of protein phosphatase 1, which in turn regulates the phosphorylation state of several classes of effector proteins, including transcription factors, ionotropic receptors, and ion channels (Greengard et al. 1999; Grace 2002). It has been shown that DARPP-32 is also present in other non-D1-containing neurons as well, including the enkephalin-containing striatal neurons (Langley et al. 1997). In this case, in contrast, stimulation of dopamine D2 receptors causes a dephosphorylation of DARPP-32 via calcineurin activation by calcium influx (Grace 2002).

Dopamine release appears to occur via two functionally distinct components. One is known as phasic component of dopamine release and is believed to underlie most of the behavioral indices of this transmitter. Phasic dopamine release occurs in a high-amplitude, brief pulsatile manner by means of action potentials and then is rapidly removed from the synaptic cleft via reuptake (Grace 1991, 2002). The other component is known as tonic dopamine release that occurs in very low concentration but is sufficient to activate extrasynaptic receptors, including dopamine terminal autoreceptors. D2 and D3 receptors act as inhibitory presynaptic autoreceptors and as postsynaptic receptors. In addition, D2 receptors have two splice variants,  $D2_{short}$  and  $D2_{long}$  (Nestler et al. 2009). Autoreceptors can exist on most portions of dopamine cells including the soma, dendrites, and nerve terminals. Stimulation of dopamine autoreceptors are supposed to exert a tonic downregulation of dopamine neuron activity, maintaining their firing within a stable range of activity (Harden and Grace 1995).

## 3 Dopamine-Related Disease: Parkinson's Disease

Several important diseases of the CNS are associated with dysfunctions of the dopamine system, ranging from PD to schizophrenia and drug addiction. In this review, we shall focus only the Parkinson's disease and some consequences for human beings. The discovery of mechanism of action of several drugs that act on dopamine synapses really improved the knowledge about the dopaminergic system and its related diseases.

PD is characterized by complex motor symptomatology, including tremor at rest, bradykinesia, rigidity, postural instability, stooped posture, and freezing that occur by hypofunctional dopamine states when nigrostriatal pathway is degenerated. Dopamine became clinically relevant when dopamine depletion in the caudate nucleus of patients with PD was discovered, whereas intravenous (i.v.) administration of L-dihydroxyphenylalanine (L-DOPA), the amino acid precursor of dopamine,

dramatically and rapidly alleviated parkinsonian symptoms (Hornykiewicz 2010). Studies with animal models of PD provided the first clear indication of the dichotomy in the regulation of the direct and indirect pathways, suggesting that their excitability shifts in opposite directions following the loss of dopamine, creating an imbalance in the regulation of the motor thalamus favoring suppression of movement (Albin et al. 1989). Specifically, direct pathway was suggested to spike less in the PD state, whereas indirect pathway was thought to spike more (Gerfen and Surmeier 2011).

Besides motor symptoms, dopaminergic-related non-motor symptoms, as depression, apathy, and sleep disorders, are part of the clinical spectrum of PD. There is increasing evidence that such symptoms are treatable, at least in part, with various dopaminergic agents (Chaudhuri and Schapira 2009).

One current drawback of dopaminergic therapy is that it cannot be targeted to a specific subpopulation of dopamine neurons. A consequence of this is significant dopamine-related side effects, such as hallucinations or compulsive gambling during treatment for PD or extrapyramidal and sexual side effects caused by antipsychotics. In addition, from the standpoint of pathogenesis, it is clear that dopaminergic neurotransmission is not the sole dysfunction in this neurological disease.

## 4 Dopamine Signaling in Parkinson's Disease

## 4.1 The Neuropathology of Parkinson's Disease: A Brief Historical Overview

The association of PD with the mesencephalic dopaminergic zone was hypothesized by Edouard Brissaud in 1893, based on an autopsy case of unilateral Parkinsonism associated with a tumor confined in the SN (see Parent and Parent 2010). The hypothesis was validated by Konstantin Tretiakoff in 1919, in his doctoral thesis, reporting the marked loss of neurons in the SNc and the presence of intracytoplasmic inclusions which he called "*corps de Lewy*" (reference to Friedrich Lewy's description of these pathological findings) in the surviving neurons of PD patients (Lees et al. 2008), after confirmed by Hassler in 1938, which found a constant cell loss in the SNc of 10 PD patients (Hassler 1938). Just on the beginning of 1960, Hornykiewicz described the presence of dopamine in SN and that it is reduced in the midbrain and in the striatum of parkinsonian patients, suggesting the possible existence of a link between SN and the corpus striatum (Hornykiewicz 1963).

Using new fluorescence techniques, Dahlström and Fuxe (1964) described firstly, in rat, the occurrence of catecholaminergic cell in the midbrain and lower brain stem, and Andén et al. (1964) noticed that lesions of SNc caused a substantial loss of the catecholaminergic fluorescence in the striatum, helping the acceptance of the existence of a nigrostriatal fiber system originating from dopaminergic midbrain neurons. The utilization of immunohistochemistry labeling of enzymes

participating in the metabolism was very useful to study the catecholaminergic neurons of midbrain, and Hökfelt et al. (1973) were the first to identify dopaminergic neurons in midbrain using antibodies to aromatic acid decarboxylase, followed by the immunohistochemical revelation of the rate-limiting enzyme of dopamine synthesis, the TH, done by Ljungdahl et al. (1975), confirming the dopaminergic nature of nigrostriatal pathway and starting a new era for basal ganglia research.

Jakes and co-workers described two proteins of synuclein family, the alpha- and beta-synuclein, in human brain (Jakes et al. 1994). After some descriptions of familial cases of PD associated with mutation on alpha-synuclein (AS) (Polymeropoulos et al. 1997; Goedert 1997), Spillantini et al. (1997) reported, for the first time, the strong immunolabeling of Lewy bodies (LB) and neurites (LN) with AS in PD and dementia with Lewy body (DLB), followed by several publications which reinforced the relevance of AS on PD pathogenesis.

## 4.2 The Topographic Braak Staging of Parkinson's Disease

After the recent observations on relationship of AS and PD, Del Tredici et al. (2002) evaluated the presence of AS in brain stem of subjects without history of neurologic or psychiatric disease. They described that all patients with AS inclusions in dorsal motor nucleus of vagus (DMV) and in intermediate reticular zone (IRtZ) presented also AS-immunoreactivity in other brain stem nuclei, as the locus coeruleus (LC) and the SN, suggesting that the pathological progression of PD could begin in the medulla and spread to the whole brain. The absence of clinical signs of PD in these individuals, even with AS pathology, can be explained by the existence of a presymptomatic phase before the onset of motor dysfunction.

The study of Braak et al. (2003) changed the classic concepts of an essentially dopaminergic dysfunction in PD. Using the same methods of Del Tredici et al. (2002) in 3 groups of (1) 41 individuals with clinical diagnosis of PD, (2) 69 individuals without neurological or psychiatric disease but with AS brain pathology, and (3) 58 individuals without neurological or psychiatric disease nor AS pathology, they showed that LB and LN marked with AS were present in nigral and extranigral areas, with a particular vulnerability and independent of neurotransmitter produced by the neurons, which became involved at different times in the progression of the disease, in a pattern of topographical caudorostral progression, as follows:

• Stage 1: The presence of LN and LB was detected in two major nuclei: DMV and the IRtZ. At this stage LN was observed first than LB; however, in advanced cases, the LN outnumbers the LB in these areas. This pattern was seen at all stages of disease progression. Some neuromelanin-laden nerve cells in these nuclei do not show LB/LN. The anterior olfactory structures (olfactory bulb/ stalk and the anterior olfactory nucleus) are also affected by AS inclusions. This stage of pathology spares the pontine melanized neurons, as the LC.

- Stage 2: The more accentuated presence of LB/LN in DMV and IRtZ and the loss of neurons are seen at this stage. Furthermore, the pathological LB/LN appears in pontine areas of caudal raphe nuclei, the gigantocellular reticular nucleus and the coeruleus-subcoeruleus complex, but the latter nucleus has unaffected neurons. The midbrain nuclei remain uninvolved.
- Stage 3: There is worsening of lesions on affected nuclei in medulla and pons, and the LB/LN is seen in midbrain structures and nuclei from the basal forebrain. In the midbrain, the presence of LB/LN in SN is observed initially in melanized projection neurons of posterolateral and posteromedial subnuclei, without macroscopically detectable depigmentation. The compact portion of pedunculopontine tegmental nucleus is affected only by LN in stage 3. The magnocellular nuclei of the basal forebrain (medial septal nucleus, interstitial nucleus of diagonal band, basal nucleus of Meynert) develop only a network of LN. There is no pathological alteration in temporal mesocortex or neocortex, but LNs are seen in anterior olfactory nucleus and second sector of Ammon's horn (CA2) (for a wide review about extranigral brain stem nuclei affected by AS pathology, see Grinberg et al. 2010).
- Stage 4: The midbrain shows accentuated involvement of neuromelanin-laden neurons of SNc, more significant in posterior region of pars compacta; the depigmentation is macroscopically seen at this stage and is observed in the extracellular melanin granules in SNc. Other mesencephalic regions are affected, such as the paranigral and parabrachial nucleus, and the compact portion of pedunculopontine tegmental nucleus (PPN) is more affected by LB and LN. The basal forebrain becomes severely involved, and areas as stria terminalis, central nucleus of amygdala, and ventral claustrum show LB. The anterior olfactory nucleus is much damaged. This stage is characterized by AS accumulation in the anteromedial temporal mesocortex, with outer layers developing networks of LN and LB within inner layers.
- Stage 5: The last two stages indicate diffuse AS immunolabeling, including the neocortex. The SN is almost without melanin, and LB develops in melanized nerve cells as DMV and reticular formation. The network of LN in CA2 extends into adjacent sectors, comprising large portions of hippocampus. From the anteromedial temporal mesocortex (the periallocortical transentorhinal region and the proneocortical entorhinal area), the pathological progression moves toward the sensory association areas, anterior cingulate cortex, and prefrontal area, with LB in infragranular layers and LN in supragranular layers. The primary neocortical areas are spared.
- Stage 6: The final stage encloses the entire neocortex, but the pathological changes on premotor area, primary motor field, and primary sensory area are mild.

The authors of this study suggested that this staging model permits the distinction of clinically manifested cases, even as non-motor initial PD patients, as REM sleep disorders, psychiatric diseases, intestinal constipation, olfactory dysfunctions, and dysautonomia. The Braak hypothesis shows a correlation among pathological and clinical phenotypes. Jellinger (2003) observed that some individuals from Braak's study had stages 3–4 PD pathology, but no motor symptoms were diagnosed, in which he called incident Lewy body pathology (iLBP). Furthermore, as noticed by Parkkinen et al. (2003) and Ishizawa et al. 2003, other diseases as DLB and AD showed LB with AS-immunoreactivity in the brain, reducing the specificity of Braak scheme for PD. After the publication of those works, the Braak staging model had gained acceptance rapidly. However, there are no conclusive proofs confirming a caudal-to-rostral temporal progression in PD pathology. Additional studies are required to elucidate these questions.

## 4.3 The Human Midbrain Dopaminergic System and Parkinson's Disease

Classically, the bradykinesia, rigidity, and tremor in parkinsonian syndromes are attributed to mesencephalic dysfunctions, and diseases as strokes and tumors localized in SN can cause a rigid-akinetic phenotype (Garcia de Yebenes et al. 1982; Akyol et al. 2006; Orta Daniel and Ulises 2008). PD is modernly considered as a multisystem neurodegenerative illness, affecting dopaminergic and non-dopaminergic neurons (serotoninergic, glutamatergic, cholinergic, nitrergic) as seen on Braak hypothesis. However, the major motor symptoms (bradykinesia and rigidity) and the motor complications (dyskinesia, wearing off phenomenon) are associated to dopaminergic transmission, justifying the special attention given to the nigrostriatal degeneration. There are other dopamine-rich areas in CNS, as the hypothalamus and retina, but its involvement in PD is controversial (for review see Archibald et al. 2009).

Based on the immunoreactivity to calbindin, the midbrain dopaminergic system can be divided into two tiers: the dorsal and the ventral tier. The dorsal tier encompasses the retrorubral groups and the VTA, its neurons are calbindin-positive, and they have horizontal orientation and low expression of DAT and D2 receptors. The ventral tier is composed by SNc, with calbindin-negative and ventrally oriented neurons, showing high amounts of DAT and D2 receptors. This cytochemical phenotypic classification has a practical application: the ventral tier is particularly vulnerable to the neurodegeneration of PD but with selective sparing of the dorsal tier (Haber and Gdowski 2004). However, the neuromelanin-laden neuron loss in SNc is not necessarily uniform (Ma et al. 1996).

Fearnley and Lees (1991) studied the micro-architecture of SN and the rates of neuron loss in normal aging and parkinsonian syndromes, and there was a linear reduction of pigmented neurons in advancing aging, more affected in dorsal tier. In opposite pattern, the PD individuals had 45 % of neuron loss in the first decade of disease, being greatest in the lateral ventral tier (average loss 91 %), followed by the medial ventral tier (71 %) and the dorsal tier (56 %). These results show a specific pattern of nerve cell loss in SNc on PD (Fearnley and Lees 1991). Ma et al. (1996) performed a morphometric analysis of SNc in PD individuals, and they described a marked reduction in pigmented neuron area, perimeter, diameter, and number in PD cases.

Damier et al. (1999a, b) developed a new method to analyze the SNc, based in immunolabeling to calbindin, identifying calbindin-rich regions ("matrix") and calbindin-poor islands ("nigrosomes"), and these pocket zone patterns were recognized in all brain as five nigrosomes. The nigrosome 1, localized in the ventral third and the lateral two-thirds of the calbindin-rich neuropil, corresponds to the ventral tier (Damier et al. 1999a). In another study (Damier et al. 1999b), the same authors used this SN analysis in PD subjects, and the neuronal depletion was maximum (98 %) in nigrosome 1, corroborating the findings of Fearnley and Lees (1991). A spatiotemporal progression of cell loss on PD was suggested, beginning in nigrosome 1, after spreading to other nigrosomes and to the matrix.

Another method used for analysis of dopamine depletion in parkinsonian syndromes is the radionuclide imaging with positive emission tomography (PET) and single-photon emission computed tomography (SPECT), applied as biomarkers for early disease and study of disease progression, mostly the presynaptic dopamine function, of the nigrostriatal pathway. There are four major ways to examine the dopaminergic terminals function: (1) the <sup>18</sup>F-dopa in SPECT and <sup>11</sup>C-dopa in PET, measuring the DOPA-decarboxylase activity; (2) molecules as <sup>11</sup>C or <sup>18</sup>F (for PET), or <sup>123</sup>I or <sup>99m</sup>Tc (for SPECT, as <sup>99m</sup>Tc-TRODAT-1) quantify the availability of DATs; (3) 11C-dihydrotetrabenazine or its 18F-labeled analogue measures the VMAT2 density; and (4) the ability of dopaminergic terminals to release dopamine after pharmacological stimuli (e.g., amphetamine) can be evaluated by changes in dopamine D2 receptor availability with <sup>11</sup>C-raclopride PET or <sup>123</sup>I-iodobenzamide SPECT (Brooks 2007; Stoessl et al. 2011).

It is known that <sup>18</sup>F-dopa uptake has correlation with nigral dopaminergic cell counts and dopamine concentrations in striatum of humans and nonhuman primates (Pate et al. 1993; Snow et al. 1993). In unilateral Hoehn & Yahr stage 1 PD patients, the dorsal posterior putamen <sup>18</sup>F-dopa uptake is first reduced, even contralateral to the asymptomatic limb, corresponding to the evidences that ventrolateral nigral dopaminergic projections to the dorsal putamen are more affected in the onset of clinical phase of PD (Kish et al. 1988).

The subsequent PD progression does not reduce the <sup>18</sup>F-dopa uptake in the first 2 years (Whone et al. 2003), but after that the functional loss occurs more rapidly in PD than aged controls (Brooks 2003), and the initial posterior-anterior gradient of striatal involvement is substituted by similar rates of decline in the whole striatum. Furthermore, the PET/SPECT measures show inverse correlation with severity of rigidity and bradykinesia in PD but poorly with tremor, suggesting the parkinsonian tremor may be associated with extranigral mechanisms (Vingerhoets et al. 1997; Benamer et al. 2000). Another finding is the increase of <sup>18</sup>F-dopa uptake into the internal globus pallidus in initial PD as a possible compensatory action, but this phenomenon vanishes with disease progression, simultaneously with onset of fluctuating responses to levodopa (Whone et al. 2003).

The motor complications on PD are also associated with radioimaging changes. PD patients with fluctuating responses to levodopa present 20 % reduction in <sup>18</sup>F-dopa uptake than those with sustained therapeutic responses (de la Fuente-Fernandez et al. 2000), and the <sup>11</sup>C-raclopride binding shows a mean 10 % fall in

posterior putamen of advanced PD subjects compared with early-stage patients (Torstenson et al. 1997). Nevertheless, PET studies with dyskinetic individuals show similar levels of striatal dopamine D1 and D2 receptor availability (Turjanski et al. 1997). The dementia in PD, a frequent non-motor complication of advanced stages, shows a reduction in <sup>18</sup>F-dopa uptake in cingulate and medial prefrontal areas, suggesting the involvement of the mesocortical dopaminergic pathway on cognitive decline in PD (Ito et al. 2002).

In conclusion, the major importance of midbrain dopaminergic neuronal degeneration on symptomatic progression of PD is unquestionable. However, it must be emphasized that there are several other non-dopaminergic neurons involved in its pathogenesis, as shown by Braak hypothesis, and the AS aggregation and accumulation can be the common link between these vulnerable nerve cells.

## 4.4 Dopamine-Related Treatments of Parkinson's Disease

The treatment of PD is very complex due to the progressive nature of the disease and the array of motor and non-motor features combined with early and late side effects associated with therapeutic interventions. While no treatments have yet been shown conclusively to slow the progression of the disease, a growing number of symptomatic pharmacologic therapies are available, as well as several forms of surgery and numerous non-pharmacological approaches.

Among dopaminergic drugs, L-DOPA standard and controlled-release formulations are marketed as fixed associations with a DOPA-decarboxylase inhibitor carbidopa or benzeraside. There are several dopamine agonists, with different receptor binding, pharmacokinetic profiles, and routes of administration and two types of indirect dopamine transmission enhancers, monoamine oxidase-B (MAO-B) and COMT inhibitors. Among non-dopaminergic medications, there are several anticholinergics, although amantadine is the only drug widely available for anti-glutamatergic effects. Functional neurosurgery for PD attempts to restore functional balance in basal ganglia relays. There are currently three targets: the ventral intermediate nucleus of the thalamus, the internal segment of the globus pallidus, and the subthalamic nucleus. Either CNS lesions (thalamotomy, pallidotomy, or subthalamic nucleus lesions) or implants of chronic stimulating electrodes at these sites (deep brain stimulation) can be used. Reconstructive surgery involves the implantation of human fetal mesencephalic cells (fetal transplant) or other dopamine-producing cells into the striatum. Rehabilitation for PD (speech, occupational, and physical therapies) has been applied empirically based on its accepted use in other chronic disorders.

It is important to consider that while drug treatments available for PD increased in the last 30 years, the first dopaminergic drug L-DOPA remains the more effective. Actually, the more recent drugs refined the treatment but they do not alter the clinical aspect and the evolution of the disease, and everyone can only temporary control and ameliorate the motor symptoms. The increasing evidences that cognitive dysfunctions, depression, constipation, pain, and nocturnal incapacity can play

Medication	Adverse effects	Indications and comments	
Anticholinergics Benztropine trihexyphenidyl	Dry mouth, dry eyes, constipation, hypotension, cognitive impairment, urinary retention	Useful for symptomatic control of Parkinson's disease (benefits are mild to moderate); associated with more adverse effects than other drugs	
Carbidopa/ levodopa Immediate- and sustained-release carbidopa/levodopa	Nausea, somnolence, dyskinesia, hypotension, hallucinations	Levodopa is the most effective medication and remains the primary treatment for symptomatic Parkinson's disease; no added benefit for motor complications with sustained-release versus immediate- release preparations	
COMT inhibitors	Diarrhea; exacerbates levodopa Useful for managing motor		
Entacapone	adverse effects; bright orange urine fluctuations ("weat	fluctuations ("wearing off" effect) in	
Tolcapone	Diarrhea; exacerbates levodopa adverse effects if dyskinesia appears; rare liver failure (liver function monitoring needed)	patients taking levodopa; levodopa dose may need to be reduced	
Dopamine agonists	Nausea, headache, dizziness	Useful for early and advanced disease	
Bromocriptine	Somnolence; hallucinations; nausea;	Useful for the initial treatment of parkinsonism and as adjunct therapy in patients taking levodopa	
Pergolide	edema; fibrosis of cardiac valves, lung, and retroperitoneum; retroperitoneal and pulmonary fibrosis		
Pramipexole	Nausea, sleep attacks, edema,	Useful for early disease and in	
Ropinirole	hallucinations, hypotension	patients with Parkinson's disease and motor fluctuations	
MAO-B inhibitors	Nausea, insomnia, drug interactions	Useful for symptomatic control of Parkinson's disease (benefits are	
Selegiline	with other MAO inhibitors/tyramine		
Rasagaline	Weight loss, hypotension, dry mouth, drug interactions with other MAO inhibitors/tyramine	mild to moderate) and as adjuvant therapy for patients with Parkinson's disease and motor fluctuations	
NMDA receptor inhibitor	Nausea, hypotension, hallucinations, confusion, edema	Useful for treating akinesia, rigidity, tremor, dyskinesia	
Amantadine			

 Table 1
 US Food and Drug Administration (FDA) approved medications for Parkinson's disease

Abbreviations: *FDA* US Food and Drug Administration, *COMT* catechol-O-methyltransferase, *MAO-B* monoamine oxidase-B, *NMDA N*-methyl-d-aspartate. Based on information from Goetz et al. (2005)

important role in the quality of life of patients and the benefits of any new treatment should be evaluated in their impact in the well-bearing, mental health, and others deficiencies of the patients. A summarized review of the pharmacological approaches currently available for PD is illustrated in Table 1.

All the dopaminergic therapies are based in the pathway of catecholamine synthesis in the body, as presented in Fig. 1. Since most parkinsonian motor

symptoms occur as a consequence of dopamine depletion in the basal ganglia, dopamine replacement strategies represent the main therapeutic approach used to counteract PD motor impairment.

## 4.4.1 3,4-Dihydroxyphenylalanine (L-DOPA)

Since its introduction in the routine treatment of PD at the beginning of the 1970s (Cotzias 1971), L-DOPA (L-3,4-dihydroxyphenylalanine), a dopamine metabolic precursor, still remains as the most effective treatment for the treatment of the classical motor parkinsonian signs. L-DOPA, the naturally occurring isomer of the amino acid 3,4-dihydroxyphenylalanine, was first isolated in 1913 from legumes (seedlings of Vicia faba) by Marcus Guggenheim (Hauser 2009). Already 2 years earlier, Casimir Funk had synthesized D.L-DOPA in the laboratory (Hauser 2009). Funk and Guggenheim considered the amino acid as a possible parent compound of adrenaline. The discovery by Peter Holtz (1959) of an enzyme, DOPA decarboxylase, in mammalian tissue (kidney) extracts that converted L-DOPA to the corresponding - biologically active - amine, that is, dopamine, represented a turning point in catecholamine research (Hauser 2009). After the discovery of dopamine occurrence in the body, many biochemical studies were performed in an attempt to provide experimental support for the proposed biosynthesis of the catecholamines, mainly dopamine, from the amino acid L-DOPA (Hornykiewicz 2002).

The introduction of high dosage of L-DOPA by George Cotzias revolutionized the treatment of PD (Singh et al. 2007). As a prodrug of dopamine, L-DOPA crosses the blood-brain barrier, and it is decarboxylated to dopamine in the nigrostriatal pathways by brain enzymes Nowadays, L-DOPA is always given with carbidopa or benserazide, peripheral DOPA-decarboxylase inhibitors, which prevents peripheral metabolism of L-DOPA and allows a higher percentage of a dose to cross the blood-brain barrier. Primarily, they block L-DOPA metabolism in the periphery, thereby reducing the rate of the first-pass metabolism, and slowing the plasma clearance of L-DOPA; however, the enzyme inhibitors do not cross the blood-brain barrier.

The effects of the dopamine replacement therapy are predictable (as are the side effects), and none of the more recently introduced synthetic dopamine agonists has surpassed the clinical benefit derived from L-DOPA. L-DOPA has a long-duration response in early disease that enables adequate symptomatic control with dosage schedules (Table 1). This is a consequence of the ability of the nigrostriatal system to convert L-DOPA to dopamine, store it in presynaptic vesicles, and release it in response to physiological stimuli. However, as the disease progresses, the conversion of L-DOPA to dopamine in dopaminergic neurons becomes limited.

## 4.4.2 L-DOPA and Dyskinesia

Although the precise mechanisms of dyskinesia have remained elusive, three main risk factors for this side effect have been conclusively identified in clinical studies, namely, a young age at disease onset, disease severity (reflecting the extent of putaminal dopamine denervation), and high doses of L-DOPA (Jenner 2008; Stocchi 2009; see a recent review of Huot et al. 2013). These risk factors are

reproduced in both nonhuman primate and rodent species by applying neurotoxic lesions to the nigrostriatal dopamine pathway, followed by daily administration of L-DOPA at a sufficient dosage.

Although initially effective, dopaminergic therapies are eventually complicated by motor fluctuations, including off-time (periods of return of PD symptoms when medication effect wears off) and dyskinesia (drug-induced involuntary movements including chorea and dystonia) in most patients (Jenner 2008; Stocchi 2009). The mechanisms by which dyskinesia develop are not completely understood, but pulsatile stimulation of dopamine receptors by short-acting agents as L-DOPA and the degree of striatal denervation have been implicated (Obeso et al. 2000; Stocchi 2009). Dyskinesia may occur at the time of maximal clinical benefit and peak concentration of L-DOPA (peak-dose dyskinesia) or may appear at the beginning and/or at the end of the L-DOPA effect (diphasic dyskinesia, off-dystonia, off-dyskinesia). The progressive reduction in the duration of the clinical effect is called "wearing off." This change in the response to L-DOPA is more related to the disease progression and loss of buffering capacity of the presynaptic terminals to keep constant the levels of dopamine in the striatum (Chase 1998). When wearing off begins, it required modification of dosage and/or dose frequency of L-DOPA or the introduction of additional therapies. Interestingly, at this stage of the disease, so long as the plasma L-DOPA concentration is kept constant, the clinical response will persist (Mouradian et al. 1990; Schapira et al. 2009) and "wearing off" does not occur if the drug is given by continuous infusion (Nutt et al. 1989; Schapira et al. 2009). Motor complications can be an important source of disability for some patients who cycle between "on" periods, which are complicated by dyskinesia and "off" periods in which they suffer severe motor disability (Schapira et al. 2009).

Nonhuman primates have been used to produce models of L-DOPA-induced dyskinesia since the late 1980s (Crossman 1987). Following 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) lesion and subsequent treatment with L-DOPA, nonhuman primates (and macaques in particular) exhibit choreiform and dystonic movements that look very similar to those seen in dyskinetic PD patients. These models have been extensively used to define the basic pharmacological features of L-DOPA-induced dyskinesia (Jenner 2003; Cenci et al. 2009). Rats with 6-hydroxydopamine (6-OHDA) lesions have been the most widely used animal model of PD since the 1970s, but the idea that these animals can exhibit L-DOPA-induced dyskinesia has become accepted only recently (Henry et al. 1998; Cenci et al. 2009). For many years, drug-induced rotation remained the only available method to study motor effects of dopaminergic therapies in this animal model. When given to unilaterally 6-OHDA-lesioned rats, L-DOPA or dopamine receptor agonists induce the animals to turn in the direction contralateral to the lesion, a behavior traditionally regarded as a correlate of the antiparkinsonian activity of these drugs. Later on, it was realized that contralateral rotation shows sensitization during chronic dopaminergic treatment (Henry et al. 1998) – a feature reminiscent of the priming for dyskinesia described in nonhuman primates and supposed to occur in PD patients. About 10 years ago came the first reports that 6-OHDA-lesioned rats treated with L-DOPA exhibit movements with dystonic and

hyperkinetic features engaging the forelimb contralateral to the lesion, as well as axial and orofacial muscles (Lundblad et al. 2002; Winkler et al. 2002). These movements interfere with the rat's physiological motor activities, and they are reduced by treatments that have antidyskinetic efficacy in nonhuman primates and PD patients. Rating scales for rodent abnormal involuntary movements (AIMs) were developed to reflect the topographic distribution, frequency, duration, and amplitude of dyskinetic behaviors induced by L-DOPA (Dekundy et al. 2007; Cenci et al. 2009). Producing and validating an animal model is a fundamental task for investigators involved in translational PD research.

Nevertheless, it must be emphasized that L-DOPA remains the "gold standard" medication in the treatment of PD. The use of other dopaminergic drugs at the beginning of the treatment helps to delay the onset of the motor complications in patients with PD. However, at some time when the disability of the patient increases, L-DOPA needs to be introduced (Olanow et al. 2009b). L-DOPA use is appropriate when all other reasonable strategies fail to produce adequate control of symptoms or when patient's symptoms pose an imminent danger to them (e.g., loss of postural reflexes with high risk for falling).

Recently, there was a growing discussion about a possible neurotoxic effect of L-DOPA that could increase the disease progression, and many patients and doctors became afraid to use the drug and delayed its use at the most (Kurlan 2005). However, a critical evaluation of older and most recent studies fails to describe a deleterious effect of L-DOPA in the evolution of the degenerative process (Olanow et al. 2004). In contrast our work showed that chronic L-DOPA treatment to partial 6-OHDA-lesioned rats induces partial recovery of the dopaminergic system suggesting their trophic effect which was confirmed by the increase of the trophic factor pleiotrophin (Murer et al. 1998; Ferrario et al. 2004).

Therefore, in the initial stages of the disease, L-DOPA therapy is the most effective for improving motor symptoms in PD. However, long-term treatment is accompanied by fluctuations in motor performance, dyskinesia, and neuropsychiatric complications. Furthermore, as PD progresses, patients develop features that do not respond well to L-DOPA therapy, such as freezing episodes, autonomic dysfunction, falling, and dementia. The increasingly diverse possibilities in the therapy of PD, and the many side effects and complications of therapy, require the formulation of reliable standards for patient care that are based on current scientific knowledge.

#### 4.4.3 Catechol-O-Methyltransferase Inhibitors (COMT)

L-DOPA is usually combined with a DOPA-decarboxylase inhibitor (carbidopa or benserazide) to improve absorption and reduce peripheral metabolism (Schapira et al. 2009). However, the majority of L-DOPA is still metabolized in the gut and liver by COMT to form 3-*O*-methyldopa (3-OMD). 3-OMD accumulates and may interfere with the brain penetration of L-DOPA. COMT inhibitors reduce the metabolism of L-DOPA, extending its plasma half-life by 30 % and prolonging the clinical effect of each L-DOPA dose. This effect was suspected to be of enough magnitude to produce more stable L-DOPA plasma levels if a COMT inhibitor was

used with each dose of L-DOPA. The hypothesis was that this combined therapy could delay the appearance of motor complications because of producing more continuous dopaminergic stimulation. However, clinical studies were unable to show this positive effect, and in opposition, the combined therapy led to earlier appearance of dyskinesia (Klivenyi and Vecsei 2010). This unexpected effect may be explained by the potentializing effect of the COMT inhibitors in the L-DOPA effect, since its dosage, or effect, is associated with earlier beginning of motor complications.

Entacapone and tolcapone are the two selective COMT inhibitors currently in use for the treatment of PD (see Table 1), while newer molecules are now in clinical evaluation. Both COMT inhibitors only have observable clinical effects when used in association to L-DOPA. The most significant clinical effect is the reduction in the daily off-time, by small prolongation in the time of L-DOPA effect (Olanow et al. 2004; Jankovic and Stacy 2007). The COMT inhibitors also potentialize the effect of L-DOPA, and they can be used to achieve a stronger symptomatic effect (Gallagher and Schrag 2008). Tolcapone is a central and peripheral COMT inhibitor and is apparently more efficacious than entacapone that only inhibits peripheral metabolism, but there are some concerns about its safety and in particular to its potential hepatotoxicity (Assal et al. 1998). Entacapone is much more widely used and appears to be less hepatotoxic (Gershanik et al. 2003; Schrag 2005). The main adverse effects are related to its potentializing dopaminergic effects and include dyskinesia, orthostatic hypotension, nausea, vomiting, and confusion. Other important side effects are the occurrence of diarrhea and urinary discoloration. The control of hepatic enzymes is always indicated during the use of tolcapone and eventually during the use of entacapone. The association of entacapone in a single tablet with L-DOPA and carbidopa is now available. Another effect of the COMT inhibitor is the reduction in the systemic levels of 3-OMD that was supposed to interfere with L-DOPA metabolism in the central nervous system. However, any clinical benefit could be proved to be due to this property (Muller and Kuhn 2006). Another systemic effect of the COMT inhibitors is the reduction in the blood levels of homocysteine in treated PD patients (Muller and Kuhn 2006; Muller and Muhlack 2009). This could be a hypothetical benefit in the long term, since homocysteine could mediate the progression of neuronal degeneration and increase the risk for development of dementia, vascular disease, and polyneuropathy in levodopa-treated PD patients (Muller and Muhlack 2009).

Recently, it was established that a continuous duodenal L-DOPA infusion may be an efficient strategy to obtain continuous dopaminergic stimulation and to decrease the complications of the therapy (Clarke et al. 2009). Other ways to obtain a stable plasma level of L-DOPA were under research and include the development of new controlled-release tablets and cutaneous patches of the drug.

#### 4.4.4 Monoamine Oxidase Isoenzyme Type B (MAO-B) Inhibitor

Substantial experimental evidences indicate that the oxidative stress is an important pathogenic mechanism in PD (Yacoubian and Standaert 2009). There are many possible mechanisms associated to the generation of free radicals in PD. The own

dopamine metabolism produces free radicals in excess. Blocking the first step of dopamine metabolism by inactivating MAO-B reduces the production of free radicals. Selegiline and rasagiline are selective MAO-B inhibitors currently in clinical use (see Table 1). MAO-B inhibition may increase the concentration of dopamine in the synaptic cleft due to the inhibition in the dopamine degradations and also produce an antiparkinsonian effect (Wu et al. 2000). Unlike selegiline, rasagiline is not metabolized to amphetamine and has no sympathomimetic activity. Rasagiline has also shown neuroprotective effects in in vitro and in vivo models of PD (Jenner 2004).

Selective MAO inhibitors become nonselective at doses greater than those recommended. In this condition, selegiline and rasagiline may increase the risk of a hypertensive crisis if used in combination with tyramine-containing foods (e.g., fermented cheese, aged meats) or amine-containing medications (e.g., over-the-counter cough/cold medicines). Serotoninergic crisis also has been associated with the use of MAO-B inhibitors in association with antidepressive drugs, since they can increase brain serotonin levels. The motor symptomatic effect of the MAO-B inhibitors is only modest, and these drugs are most commonly used to treat patients at the earlier stage of the disease. However, the MAO-B inhibitors may also be used at the later stages to increase the daily on-time. Despite recent clinical studies with rasagiline suggested that the drug may also have a neuroprotective effect (Olanow et al. 2009a), this is still a matter of controversy (see Ahlskog and Uitti 2010).

#### 4.4.5 Dopamine Receptor Agonists

Dopamine agonists directly stimulate dopamine receptors. Of the nine dopamine agonists marketed for the treatment of PD, five were ergot derivatives (bromocriptine, cabergoline, dihydroergocryptine, lisuride, and pergolide) and four are non-ergot derivatives (apomorphine, piribedil, pramipexole, and ropinirole) (see Table 1). Pergolide was retired of the market of most countries due to tendency to induce cardiac valvar fibrosis; however, the use of ergot-derived dopamine agonists (pergolide and cabergoline) in patients with PD was associated with increased risk of cardiac valvar regurgitation (Steiger et al. 2009). The acute side effects of dopamine agonists are similar to those observed with L-DOPA and include nausea, vomiting, and postural hypotension. It is generally accepted that the shared D2-like receptor agonistic activity produces the antiparkinsonian effect. This D2 effect also explains peripheral (gastrointestinal nausea and vomiting), cardiovascular (orthostatic hypotension), and neuropsychiatric (somnolence, psychosis, and hallucinations) side effects. Even at lower doses, patients experience orthostatic hypotension, constipation, dyskinesia, confusion, and excessive somnolence with these agents (Wong et al. 2003). These adverse effects tend to occur with the initiation of treatment and to abate as tolerance develops usually over the ensuing days to weeks. Other problems related to the use of dopamine agonists include weight gain (possibly related to overeating) and edema (especially in the lower extremities). More recently, attention has been focused on the association of dopamine agonist treatment with the development of pathologic gambling, hypersexuality, and compulsive eating and shopping. This has attracted considerable scientific interest because of the known relationship between dopamine and reward (Obeso et al. 2000).

Almost all currently used dopamine agonists are able to protect dopamine neurons from death in many in vitro and in vivo experiments. This neuroprotective effect may be the result of different mechanisms: antioxidation, scavenging of free radicals, suppression of lipid peroxidation, and inhibition of apoptosis. Dopamine agonists have a significant clinical effect in the motor parkinsonian signs, but most of them were not comparable to that observed with L-DOPA. The use of dopamine agonists at the beginning of the treatment of PD to spare L-DOPA is indicated specially for younger patients. The monotherapy with this class of drugs is associated with lower incidence of dyskinesia and wearing off but with a lower antiparkinsonian effect than L-DOPA (Hitzeman and Rafii 2009). One explanation for their lower risk to induce motor complications may be due to the long half-life of these drugs in relation to the L-DOPA. One exception for this is apomorphine that has a pharmacokinetic curve very similar to the L-DOPA and is the only dopamine agonist which a similar symptomatic effect to it. The main limitation for the clinical use of this drug is the administration route that must be subcutaneous. In some countries apomorphine is infused subcutaneously and continuously by a pump mechanism in patients with PD. The clinical effect is significant and persistent (Clarke et al. 2009). This is now recognized as a potential continuous dopaminergic stimulation strategy and reduces functional impairment and motor complications.

New delivery ways for dopamine agonist molecule are in development. Subcutaneous patches of rotigotine seem to be efficient and may produce more continuous dopaminergic stimulation (Chen et al. 2009). Long-release tablets of non-ergot dopamine agonists are now in evaluation and may be useful to increase adherence of the patient to the therapeutics.

## 4.4.6 Is There Nitric Oxide Inhibition Utility in the Dopamine Replacement Therapy?

Notwithstanding recent advances in the management of PD, L-DOPA-induced dyskinesia continues to be a clinical and therapeutic challenge (Cenci and Lindgren 2007; Jenner 2008; Santini et al. 2008). However, many other neurotransmitters may play a central role in the neurobiology and management of L-DOPA-induced dyskinesia. Nitric oxide is a free radical that can also act as an atypical neurotransmitter and influence dopamine-mediated neurotransmission. Nitric oxide also regulates synaptic and neural plasticity being enrolled in the regulation of motor activity (Pierucci et al. 2011; Del Bel et al. 2005, 2011). There is evidence whereby communication between nitrergic and dopaminergic systems plays an essential role in the control of the nigrostriatal pathway. The dual localization of immunoreactivity for nitric oxide synthase (NOS) and tyrosine hydroxylase, enzymes responsible for the synthesis of nitric oxide and dopamine, respectively, proposes a close anatomical link between the two neurotransmitters (Mitkovski et al. 2012).

Recently, our group (Padovan-Neto et al. 2009, 2011; Novaretti et al. 2010; Del Bel et al. 2011) and other laboratories (Takuma et al. 2012; Yuste et al. 2011)

described that nitric oxide synthase inhibitors are able to decrease the motor changes related to prolonged administration of L-DOPA in rodent models of PD. Similar results were described, in latest meeting communication, for nonhuman primates by Iravani et al. (2008) and Yuste et al. (2011). Postmortem studies in human brain of patients with Parkinson's disease have greatly contributed to figure out the disease. In spite of this, few human brain studies have focused on the relationship between the development of dyskinesia and biochemical changes in the human brain. It limits the possible understanding of the nitric oxide system in the treatment of Parkinson's disease. But Parkinsonian patients presented in serum a marked rise of the production of the nitric oxide second messenger cvclic guanosine monophosphate (cGMP) levels (Chalimoniuk et al. 2004) induced by L-DOPA. Similar result was described in the cerebellum (Gumulka et al. 1976). Comparable result was described in animals (Itokawa et al. 2006). In mice, cGMP is regulated by L-DOPA (Chalimoniuk and Langfort 2007). In addition, Sanchez and cols. (2002) showed that nitric oxide stimulates L-DOPA release in the striatum in a time- and concentration-dependent manner.

A role for nitrergic neurotransmission in the pathophysiology of dyskinesia and as a potential drug target for new therapeutic has been proposed also by others (Iravani and Jenner 2011; Pierucci et al. 2011; Huot et al. 2013). NOS inhibitor treatment was able to prevent abnormal involuntary movements without significantly changing normal motor (Padovan-Neto et al. 2009). Different from the cataleptic effects of these drugs, there was no tolerance to the antidyskinetic effects of NOS inhibitors (Novaretti et al. 2010). The antidyskinetic effects of 7-nitroindazole, a preferential neuronal nitric oxide synthase (nNOS) inhibitor, seem to involve reduction in gene and protein expression in denervated striatal neurons (Padovan-Neto et al. 2011). L-DOPA increased nNOS mRNA levels in the contra- and ipsilateral side to the dopamine lesion of the frontal cortex but did not produce any further increases of nNOS protein in the striatum compared to the 6-OHDA-induced increase (Padovan-Neto et al. 2011).

Despite promising studies, efforts in synthesizing highly selective nNOS inhibitors have been difficult and have limited the development of such drugs (Hobbs et al. 1999; Vallance and Leiper 2002). To date, clinical studies of NOS inhibitors have been limited by the small number of potent and selective drugs that are safe for human administration (Vallance 2003; Petros et al. 1994: Rees 1995; Ashina et al. 1999). The recent discovery that multiple subunits of NOS exist, with apparent unique anatomic distribution and functional and pharmacological diversity (for review see Garthwaite 2008), holds considerable therapeutic potential. One novel therapeutic approach could be to manipulate the levels of endogenous NOS inhibitors. NOS inhibitors are also likely synthesized within the body. L-NMMA (NG-monomethylarginine) and asymmetric dimethylarginine are synthesized by methylation of arginine residues in proteins (Leiper and Vallance 1999). Once the proteins are broken down, the free methylarginines are released into the cell cytosol and act as competitive inhibitors of all NOS isoforms. They are metabolized to citrulline by the action of dimethylarginine dimethylaminohydrolase (Leiper and Vallance 1999; MacAllister et al. 1996). Inhibition of this enzyme leads to the accumulation of methylarginines and thereby blocks nitric oxide production. Antagonists might be developed to target-specific pathways within the basal ganglia.

A novel class of drugs named membrane-associated guanylate kinase inhibitors has recently been proposed in treatment of excitotoxicity (Lau and Tymianski 2010; Doucet et al. 2012). These peptides are competitive antagonists designed to bind postsynaptic scaffolding proteins (as, e.g., PSD-95, postsynaptic density 95 kDa), aimed at dissociating the spatial relationship between nNOS and NMDA (*N*-methyl-d-aspartate) receptors (NMDA/PSD-95/nNOS receptor complex) (Zhou et al. 2010; Doucet et al. 2012). The association of nNOS with NMDARs, via PSD-95 (Christopherson et al. 1999), plays an important role in a range of normal neuronal functions including synaptic plasticity (Garthwaite et al. 1988; Steinert et al. 2010) as well as pathophysiological disorders of the brain (Aarts et al. 2002; Sun et al. 2008; Zhou et al. 2010; Doucet et al. 2012). Since (a) NMDARs couple to nNOS and (b) NMDAR antagonists (amantadine) and nNOS inhibitors possess antidyskinetic properties, the disruption of the NMDAR/PSD-95/nNOS complex may possibly represent an alternative approach to specifically block nitrergic signaling coupled to NMDA receptor activity. It may prevent the problems associated general reduction of NMDA or nNOS function. To date, PDZ-based interactions represent a drug target that still remains largely unused (Dev 2004). They are however currently undergoing phase II clinical trials (Lau and Tymianski 2010).

The reduction in L-DOPA-induced dyskinesia in rodents and primates treated with NOS inhibitors suggests the involvement of nitric oxide in this altered behavior. These preclinical findings suggest that nitric oxide is a promising therapeutic target for the reduction of L-DOPA-induced dyskinesia.

## 5 Conclusion

The discoveries of dopamine as a neurotransmitter in the brain, its depletion in patients with PD, and its replacement with L-DOPA therapy were major revolutionary events in the rise to effective therapy for patients with PD. However, although drugs that act on the dopaminergic system provide effective symptomatic control for the classic parkinsonian motor features, the patients' quality of life continues to deteriorate as a consequence of gait and balance difficulties, autonomic dysfunction, and cognitive impairment. L-DOPA remains at present the most powerful symptomatic drug for the treatment of this condition. However, motor complications of chronic L-DOPA treatment have emerged as a major limitation of this therapy. Nonetheless, the importance of dopamine neurotransmitter system in a broad array of human disorders ranging from PD to schizophrenia has driven an intensive array of investigations oriented toward increasing our understanding of this complex system in normal conditions as well as disease states. Knowledge of dopaminergic cells and circuits has been enriched by findings obtained with several and highly sensitive techniques, as immunohistochemistry, cellular biology, transgenic mice or conditional mutants, functional anatomy, electrophysiological studies, sophisticated behavioral analyses, and functional neuroimaging. Altogether, these studies have also led to new concepts of healthy and pathological functions of dopaminergic circuits at the molecular and system levels.

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# Iron Neurotoxicity in Parkinson's Disease

# Marco T. Núñez, Pamela Urrutia, Natalia Mena, and Pabla Aguirre

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M.T. Núñez (🖂) • P. Urrutia • N. Mena • P. Aguirre

Department of Biology, Faculty of Sciences, University of Chile, Casilla, Santiago, Chile e-mail: mnunez@uchile.cl

#### Abstract

Iron plays essential roles in the development of early cognitive functions and in the maintenance of neuronal functions in the mature brain, so neurons have in place expeditious mechanisms to ensure a readily available supply of iron. However, iron is an intrinsic producer of reactive oxygen species (ROS), and increased levels of iron promote neurotoxicity because of hydroxyl radical formation, glutathione consumption, protein aggregation, lipid peroxidation, and nucleic acid modification. In this chapter, the components of iron homeostasis and the mechanisms by which iron homeostasis is lost in Parkinson's disease (PD) are discussed. In particular, it will be discussed the relevance of endogenous toxins such as aminochrome as mediators of mitochondrial dysfunction and of hepcidin as a mediator of inflammatory stimuli. A model is proposed that describes a positive feedback loop between changes in iron, glutathione, and ROS levels that ends in cell death.

#### Keywords

DMT1 • FPN1 • Glutathione • Hepcidin • Iron • Iron accumulation • Iron chelation • IRP1 • Mitochondrial complex I • Mitochondrial dysfunction • MPTP • NTBI • Parkinson's disease • Reactive oxygen species

List of Abbreviations			
AC	Aminochrome		
CDC14A	Dual specificity protein phosphatase		
CNS	Central nervous system		
CP	Ceruloplasmin		
CSF	Cerebrospinal fluid		
DAT	Dopamine transporter		
Dcytb	Duodenal cytochrome b ferrireductase		
DMT1	SLC11A2, divalent metal transporter 1		
EMSA	Electrophoretic mobility shift assay		
FPN1	SCL40A, ferroportin 1		
GPAT	Glutamyl phosphoribosyltransferase		
GSH	Glutathione		
GSSG	Oxidized glutathione		
HIF	Hypoxia-inducible factor		
IFN-γ	Interferon gamma		
iNOS	Inducible nitric oxide synthase		
IRE	Iron regulatory element		
IRP	Iron regulatory protein		
L-DOPA	L-3,4-dihydroxyphenylalanine		
LIMK1	LIM domain kinase 1		
LIP	Labile iron pool		
LPS	Lipopolysaccharide		
MPP+	1-methyl-4-phenylpyridinium		
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine		
MRCKa	Myotonic dystrophy kinase-related Cdc42-binding kinase $\alpha$		

ΝFκB	Nuclear factor kappa B
NMDA	N-methyl-D-aspartate
NO	Nitric oxide
NTBI	Non-transferrin-bound iron
PAP7	PKA-associated protein 7
PD	Parkinson's disease
PINK-1	PTEN-induced putative kinase 1
ROS	Reactive oxygen species
SDR2	Stromal cell-derived receptor 2
SN	Substantia nigra
Tf	Transferrin
TfR1	Transferrin receptor 1
TH	Tyrosine hydroxylase
TNF	Tumor necrosis factor
TNFR	TNF receptor
TRPC6	Transient receptor potential canonical 6
XO	Xanthine oxidase
Zip 14	Zrt-, Irt-like protein 14

## 1 Introduction

Early observations in postmortem PD tissue showed decreased glutathione (GSH) and increased oxygen-mediated damage as detected by 4-hydroxynonenal protein adducts (Perry et al. 1982; Youdim et al. 1989; Jenner 1991). These findings established the evidence for the participation of oxidative damage in nigral neuronal death occurring in PD. At present, the causes for this deregulation remained obscure.

An important development in the study of the molecular mechanisms underlying the disease arose from the serendipitous discovery of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) as a Parkinsonian agent (Langston et al. 1983). This initial finding was followed by the finding that the primary effect of MPTP, inhibition of mitochondria electron transport chain complex I, was a customary characteristic of Parkinsonian tissue (Schapira et al. 1990). Thus, it became fairly well established that mitochondrial dysfunction and the consequences therein are most probably germane to the development of the disease.

The finding of genetic mutations that induce familiar Parkinsonism has provided important insights into the cellular and molecular processes that go awry in PD (for a recent review see (Martin et al. 2011)). PINK1, parkin, DJ-1, and  $\alpha$ -synuclein are proteins that interact with the mitochondrion and modulate its function. The fact that their loss of function results in familiar PD supports the notion that mitochondrial dysfunction is a central event to PD pathogenesis (Schapira and Gegg 2011).

There is abundant evidence showing iron accumulation in dopaminergic neurons of the SN pars compacta (Jellinger 1999; Bartzokis et al. 2000; Sayre et al. 2000; Perry et al. 2003; Zecca et al. 2004; Berg and Youdim 2006; Wilson 2006; Weinreb et al. 2011).

The observation that pharmacological agents with iron chelation capacity prevent neuronal death induced by Parkinsonian toxins (Zhu et al. 2007) highlights the pivotal role of iron in PD neuronal death. It is therefore important to understand the relationships between mitochondrial complex I dysfunction, oxidative stress, and iron accumulation and how the conjunction of these factors results in neuronal death. The pivotal role of iron in the events leading to nigral neuronal death is the subject of this chapter.

## 2 Iron Homeostasis and Dyshomeostasis: The Role of Iron Transporters on Iron Accumulation

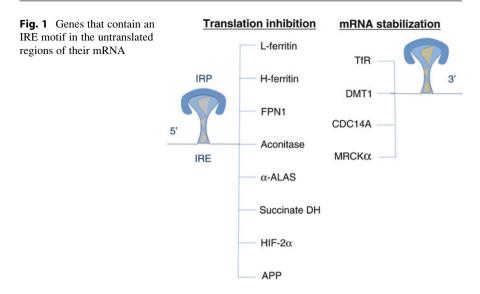
## 2.1 The Iron Responsive Element: Iron Regulatory Protein (IRE/IRP) System. A Brief Description

In vertebrates, cellular iron levels are posttranscriptionally controlled by the activity of iron regulatory proteins (IRP1 and IRP2), cytoplasmic proteins that bind to structural elements called iron-responsive elements (IREs). IREs are found in the untranslated region of the mRNAs of the major proteins that regulate cellular iron homeostasis: the transferrin receptor, the iron transporter DMT1, involved in iron import, the iron-storage protein ferritin and the exo-transporter ferroportin 1, involved in decreasing the levels of cytosolic redox-active iron (for a recent review see (Wang and Pantopoulos 2011)).

The activities of both IRP1 and IRP2 respond to changes in cellular Fe status, although through different mechanisms. In iron-replete conditions, IRP1 has a 4S-4Fe cubane structure that renders the protein active as a cytoplasmic aconitase but inactive for IRE binding. Low levels of intracellular Fe induce disassembling of the 4S-4Fe cluster, which causes IRP1 to bind and stabilize TfR and DMT1 mRNA. Besides being regulated by iron levels, IRP1 is also regulated by effectors such as nitric oxide (Bouton 1999), hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) (Martins et al. 1995), hypoxia (Hanson et al. 1999), and phosphorylation (Schalinske and Eisenstein 1996). In contrast to IRP1, IRP2 activity is downregulated through iron-induced oxidative damage followed by ubiquitination and proteasome degradation (Guo et al. 1995). The genetic ablation of IRP2 revealed that IRP2-/- mice are born normal but develop in adulthood a movement disorder characterized by ataxia, bradykinesia, and tremor (LaVaute et al. 2001). In contrast, IRP1 - / - mice are apparently normal during adulthood, with only slight misregulation of iron metabolism in the kidney and brown fat (Meyron-Holtz et al. 2004). Thus, IRP2 seems to dominate the physiological regulation of iron metabolism, whereas IRP1 seems to predominate in pathophysiological conditions.

A genome-wide search for IRE sequences resulted in 35 new target genes for IRPs (Campillos et al. 2010; Sanchez et al. 2011). Thus, the scope of the IRE/IRP system has lately gone from a restricted iron homeostasis system to a more global system, still undetermined, possibly linking iron homeostasis to a myriad of other cell functions (Fig. 1).

Newly found genes that contain an IRE motif include Hif- $2\alpha$ , which regulates the expression of proteins involved in erythropoiesis, angiogenesis, programmed

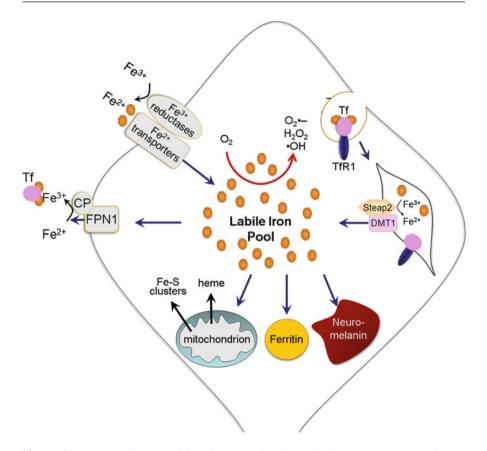


cell death, cancer, and ischemia (Patel and Simon 2008; Davis et al. 2011). Other prominent members are Cdc14, a tyrosine phosphatase involved in exit from mitosis and initiation of DNA replication. Cdc14 has been shown to dephosphorylate tumor suppressor protein p53, possibly regulating its function and MRCK $\alpha$ , a serine/threonine kinase regulated by Cdc42, a member of the Rho subfamily of small GTPases that regulate cytoskeletal organization. MRCK $\alpha$  also regulates myosin assembly by phosphorylation of the myosin-binding subunit of myosin light chain and phosphorylates LIMK1 and LIMK2, kinases that regulate actin assembly (Cmejla et al. 2010).

## 2.2 Cellular Components of Neuronal Iron Homeostasis

The components of neuronal iron homeostasis are shown in Fig. 2.

The scheme includes transferrin and transferrin receptor (TfR), inflow (DMT1; SLC11A2) and efflux (ferroportin 1, FPN1) iron transporters, the iron-storage protein ferritin, the ferrireductase Dcytb that reduces extracellular  $Fe^{3+}$  to  $Fe^{2+}$  prior to transport by DMT1, and the ferroxidase ceruloplasmin (CP) responsible for the oxidation of  $Fe^{2+}$  after transport by FPN1 and prior to the binding by apoTf. Transferrin-bound iron uptake starts with the binding of transferrin to surface receptors, followed by internalization into the endosomal system, release of iron mediated by the acidification of the endosome, reduction possibly mediated by Steap2, and transport into the cytoplasm by endosomal DMT1. Once in the cytoplasm,  $Fe^{2+}$  becomes part of the labile or reactive iron pool where it distributes to mitochondria, neuromelanin, and ferritin or engages in electron exchange reactions (Kakhlon and Cabantchik 2002; Kruszewski 2003). All the components described in Fig. 1 have been detected in the brain (Moos et al. 2007; Rouault et al. 2009; Haeger et al. 2010).



**Fig. 2** Components of neuronal iron homeostasis. The molecular components comprise the transferrin-transferrin receptor complex; the ferrireductases Dcytb and/or SDR2, responsible for the reduction of  $Fe^{3+}$  prior to transport by DMT1; uptake (DMT1, TRPC6, VGCC, Zip14) and efflux (ferroportin 1, FPN1) iron transporters; the iron-storage protein ferritin; the ferrireductase Steap2, probably responsible for the reduction of  $Fe^{3+}$  prior to transport by DMT1 in endosomes in the brain; and the ferroxidase CP, responsible for the oxidation of  $Fe^{2+}$  after transport by FPN1 and prior to  $Fe^{3+}$  binding to apoTf. Once in the cytoplasm, iron forms part of the labile iron pool from where it distributes to mitochondria, ferritin, and neuromelanin, or it is exo-transported by FPN1. In the cellular environment, LIP iron is a net ROS producer. For simplicity, the molecular components of mitochondrial iron homeostasis were omitted (From Núñez et al. (2012) with permission from the publisher)

#### 2.2.1 The Iron-Uptake Transporters

Iron concentration in cerebrospinal fluid (CSF) ranges between 0.2 and 1.1  $\mu$ M, whereas transferrin concentration lies around 0.24  $\mu$ M (Moos and Morgan 1998; Symons and Gutteridge 1998). Thus, CSF iron often exceeds the binding capacity of transferrin, and non-transferrin-bound iron (NTBI) uptake is expected to occur in neurons that express DMT1 or other iron-uptake transporters.

DMT1 (SLC11A2) is the paradigmatic transporter that brings iron inside cells (Gunshin et al. 1997). In the brain, DMT1 is expressed in hippocampal pyramidal

and granule cells, cerebella granule cells, pyramidal cells of the piriform cortex, SN and the ventral portion of the anterior olfactory nucleus, striatum, cerebellum, hippocampus and thalamus, as well as in vascular cells throughout the brain and ependymal cells in the third ventricle (Gunshin et al. 1997; Williams et al. 2000; Burdo et al. 2001). The pervasive presence of DMT1 in neurons suggests that DMT1 is necessary for their regular function (Hidalgo and Núñez 2007; Wright and Baccarelli 2007; Haeger et al. 2010; Pelizzoni et al. 2011).

The mammalian DMT1 gene undergoes alternative splicing generating four isoforms, all with similar activities for Fe<sup>2+</sup> transport. The 1A and 1B mRNA DMT1 variants originate from alternative splicing at the 5' end (exons 1A and 1B), while the +IRE or –IRE variants originate from splicing on the 3' end (exons 16/16A and 17) (Hubert and Hentze 2002; Ludwiczek et al. 2007). The 1B isoforms are expressed in the brain (Haeger et al. 2010), whereas the 1A isoforms are almost exclusively expressed in the intestine (Hubert and Hentze 2002).

Given its crucial role on cell iron uptake, knowledge on the regulation of DMT1 expression is vital to understand pathological conditions in which iron accumulation is observed. It is generally accepted that the +IRE isoforms of DMT1 are regulated by the IRE/IRP system, which posttranscriptionally regulates the expression of iron homeostasis proteins such as transferrin receptor 1 (TfR1), DMT1, FPN1, and ferritin, in response to the concentration of reactive iron in the cytoplasm (Garrick and Garrick 2009). Knowledge on the regulation of DMT1(–IRE) mRNA levels through its 3'-UTR is scarce. Recently, its regulation by the levels of miRNA Let-7d was reported in developing erythroid cells (Andolfo et al. 2010). The authors demonstrated that when the levels of miRNA Let-7d increase, the expression of DMT1(–IRE) decreases, probably by inducing its degradation through binding to imperfect complementary sites within the 3'-UTR of its mRNA.

Understanding on the transcriptional regulation of DMT1 expression is emerging. The inflammatory cytokine NF $\kappa$ B and nuclear factor Y regulate DMT1(1B) expression in embryonic carcinoma cells (Paradkar and Roth 2006b), whereas hypoxia upregulates expression of the DMT1(1A) isoform, presumably through activation of HIF-1 (Lis et al. 2005; Wang et al. 2010a). HIF-1 also promotes the expression of the 1B isoform by binding to a putative hypoxia-response element sequence in DMT1 exon1B gene (Qian et al. 2011). Recently, upregulation of DMT1 mRNA by the endogenous neurotoxin aminochrome was reported (Aguirre et al. 2012). Micromolar concentrations of aminochrome induced increased DMT1(+IRE) expression coupled to decreased FPN1 expression, in a process blocked by the antioxidant N-acetyl cysteine. Thus, it is possible that an increase in the oxidative tone triggers, through an undefined mechanism, increased expression of DMT1.

#### 2.2.2 The Iron Exo-transporter FPN1

FPN1 (SCL40A) is the only member of the SLC40 family of transporters and the first reported protein that mediates the exit of iron from cells (McKie et al. 2000). The protein is expressed mainly in enterocytes and macrophages, but its presence has also been described in neurons and astrocytes (Burdo et al. 2001).

In enterocytes, FPN1 is responsible for iron efflux during the process of intestinal iron absorption, while in Kupffer cells FPN1 mediates iron export for reutilization by the bone marrow (Devalia et al. 2002).

Knowledge on the regulation of FPN1 expression is incipient. Most probably, FPN1 expression is regulated at both transcriptional and translational levels (McKie et al. 2000; Knutson et al. 2003). There are two closely related isoforms of FPN, one (FPN1) that has an IRE motif in the 5'-UTR and another (FPN1B) that does not have it (Zhang et al. 2009). The 5'-UTR motif gives FPN1 a ferritin-like response to variations in cell iron, increasing its expression under elevated cell iron conditions (McKie et al. 2000; Yang et al. 2002; Aguirre et al. 2005). For the contrary, the 1A isoform, mainly expressed in enterocytes and erythroid precursor cells, responds to an iron challenge decreasing its expression (Zhang et al. 2009).

The expression of FPN1 in mouse brain is ubiquitous. It is present in oligodendrocytes, microglia, astrocytes, and neurons (Song et al. 2010). Space-temporal expression of FPN1 in neurons is variable (Moos and Rosengren 2006). In young brain high immunoreactivity is found in the neurons of the hippocampus and striatum (both in cell bodies and in projection fibers), a mild expression in the SN pars compacta and the superior colliculus and low expression in the SN pars reticulata (Boserup et al. 2011). In the adult brain, FPN1 immunoreactivity is lower in the projections of the striatum, but no differences have been found in neuronal cell bodies (Moos and Rosengren 2006).

## 2.2.3 TfR1

The TfR1 mediates the incorporation of iron into cells by binding the plasma irontransport protein transferrin and undergoing recycling clathrin-mediated endocytosis (Harding et al. 1983). The presence of TfR in neurons, microglia, and astrocytes and its response to iron deficiency have been described (Han et al. 2003; Beard et al. 2005). Early studies by Faucheux et al. revealed that the density of <sup>125</sup>I-labeled transferrin binding sites were highest in the central gray substance, intermediate in the catecholaminergic cells of the superior colliculus and ventral tegmental area, and almost nonexistent in the SN (Faucheux et al. 1993).

As noted above, TfR1 is posttranscriptionally regulated by the IRE/IRP system. When cellular iron is low, IRPs bind to five IRE motifs in the 3'-UTR of TfR1 mRNA increasing its mRNA stability. TfR1 expression is transcriptionally regulated by hypoxia through HIF-1 (Tacchini et al. 1999). TfR1 expression is also regulated by inflammatory stimuli: LPS/IFN- $\gamma$  treatment to mouse macrophages upregulates NF- $\kappa$ B, which in turn transiently activates HIF-1-dependent TfR1expression and iron uptake (Tacchini et al. 2008).

## 2.2.4 Ferritin

Ferritin, a multimeric protein formed by 24 subunits of H and L monomers, is the only well-characterized iron-storage protein in living organisms (Liu and Theil 2005). The 24 subunits in ferritin form a hollow cavity that can store as much as  $4,500 \text{ Fe}^{3+}$  atoms in the form of crystallized diferric oxo-hydroxyl complexes (Harrison and Arosio 1996). Ferritin plays a fundamental role in controlling the

size of the cytoplasmic redox-active iron pool (Salgado et al. 2010). Although ferritin is expected to reduce this pool by storing iron, there is a dynamic exchange of iron between its ferritin-bound form and a cytoplasmic form amenable to transport by FPN1 (De Domenico et al. 2006). In addition, ferritin contributes to the pool of redox-active iron every time it turns over (Mehlhase et al. 2005). H-ferritin, the subunit responsible for  $Fe^{2+}$  oxidation, is relatively poorly expressed in melanized dopaminergic neurons of the SN pars compacta, as compared to neurons in other parts of the brain, but is strongly expressed in oligodendrocytes (Snyder and Connor 2009).

#### 2.2.5 The Ferrireductases and Ferroxidases

 $Fe^{2+}$  is the preferred species for transport, so ferrireductases must reduce  $Fe^{3+}$  prior to its transport into the cell. In contrast, Fe<sup>2+</sup> transported from inside the cell to the extracellular milieu must be oxidized by ferroxidases to  $Fe^{3+}$  prior to transferrin binding. Three membrane ferroxidases have been described: Dcytb, its homologous cytochrome b561, and SDR2 (Vargas et al. 2003). Dcytb, initially described as the ferroxidase responsible for reduction of nonheme iron in the duodenal lumen (McKie et al. 2001), is the only member of the iron transport machinery that lacks an IRE in its mRNA. Recently, the upregulation of Dcytb by Hif- $2\alpha$  was reported (Latunde-Dada et al. 2011). As mentioned before, DMT1 and FPN1 exhibit transcriptional regulation by Hif-1. Thus, the control of iron acquisition through the upregulation of Dcytb and DMT1 by Hif-1, coupled to the downregulation of FPN1 by Hif-2 $\alpha$ , results in a concerted response of increased iron accumulation under hypoxic conditions. The presence of Dcytb and SDR2 was recently reported in astrocytes, where they may have a limited role in iron accumulation by these cells (Tulpule et al. 2010), but there is no report on their presence in neurons. Maybe the high concentration of ascorbate in the CSF, which should keep NTBI in the 2+ state, makes unnecessary a membrane-bound iron reduction system in brain cells.

The Steap family of ferrireductases comprises the members Steap1, Steap2, Steap3, and Steap4 (Ohgami et al. 2006). Steap3 was described in endosomes of erythroid precursor cells, catalyzing the reduction of  $Fe^{3+}$  to  $Fe^{2+}$  prior to transport by DMT1 (Ohgami et al. 2005). In situ hybridization studies show that Steap2 is expressed in the brain (Ohgami et al. 2006). Whether Steap2 or other member of the Steap family has a role in iron uptake by neurons remains speculative.

Two multi-copper ferroxidases, ceruloplasmin (CP) and hephaestin (Heph), are involved in iron export from the cell. CP seems to be required for iron release from most tissues in the body. Although the liver is the predominant source of serum ceruloplasmin, ceruloplasmin gene expression has also been demonstrated in many tissues including spleen, lung, testis, and brain (Hellman and Gitlin 2002). The scope of action of the membrane-bound CP homolog Heph is restricted to the small intestine, where it is required for efficient iron absorption (Vulpe et al. 1999).

#### 2.2.6 Neuromelanin

Neuromelanin is a polymer formed by oxidized metabolites of dopamine, containing a peptide component of about 15 % (Zecca et al. 2002). Neuromelanin

avidly binds iron in its +3 form, presenting high- and low-affinity binding sites (Double et al. 2003). Given the low expression of ferritin, neuromelanin is the main iron-storage moiety in SN pars compacta neurons. It is believed that the high-affinity sites are protective as they sequester iron in a redox-inactive form, whereas iron in the low-affinity sites is redox active (Gerlach et al. 2008). Thus, when iron concentration increases above the high-affinity binding capacity of neuromelanin, the neurons become prone to oxidative damage.

## 3 Iron Accumulation and Glutathione Depletion in PD

Iron is an intrinsic ROS producer. When one or more of its six ligand binding sites are not tightly bound, iron engages in one-electron exchange reactions, producing free radicals (Graf et al. 1984) through the Haber-Weiss and Fenton reactions. The thermodynamic balance of these reactions indicates that in the reductive environment of the cell, iron, in the presence of oxygen, catalyzes the consumption of GSH and the production of the hydroxyl radical ('OH) (Núñez et al. 2012).

In dopaminergic cells, another source of free radicals derives from the nonenzymatic oxidation of dopamine mediated by redox-active iron, resulting in semiquinones and  $H_2O_2$  production (Zoccarato et al. 2005). Thus, redox-active iron, both through the Fenton reaction or via dopamine oxidation, is a dangerous prooxidant agent.

## 3.1 Iron Accumulation in PD

Overwhelming evidence indicates that iron accumulation is a common feature of a number of neurodegenerative disorders of the central nervous system that include Huntington's disease, Alzheimer's disease, Friedreich's ataxia, amyotrophic lateral sclerosis, and Parkinson's disease (Jellinger 1999; Bartzokis et al. 2000; Sayre et al. 2000; Perry et al. 2003; Zecca et al. 2004; Berg and Youdim 2006; Wilson 2006; Weinreb et al. 2011). Indeed, the autocatalytic production of hydroxyl radicals because of iron accumulation has been proposed as the unifying mechanism of cell death in neurodegenerative diseases (Kell 2010).

Iron accumulation has been demonstrated in the dopaminergic neurons of the SN pars compacta (Youdim et al. 1989; Hirsch et al. 1991; Gorell et al. 1995; Vymazal et al. 1999). Interestingly, MPTP, a drug that causes experimental Parkinson's disease, upregulates DMT1(+IRE) protein expression in mice ventral mesencephalon, where it increases neuronal death presumably through abnormal increases in cellular iron content (Salazar et al. 2008; Jiang et al. 2010). Additionally, DMT1(–IRE) mediates L-DOPA neurotoxicity in primary cortical neurons (Du et al. 2009).

The contribution of iron dyshomeostasis to the progression of events leading to neuronal death is likely, since iron accumulation has been detected in brain tissue from patients who have died, after the final steps of the pathology. Since neuronal death caused by MPTP or 6-hydroxydopamine intoxication is blocked by pharmacologic or genetic chelation of iron (Kaur et al. 2003; Shachar et al. 2004; Youdim et al. 2004; Youdim and Buccafusco 2005; Zheng et al. 2010) or by dysfunction of the iron transporter DMT1 (Salazar et al. 2008), it is possible that iron dyshomeostasis takes place in the late stages of the disease as part of a vicious cycle resulting in uncontrolled oxidative damage and cell death. A recent study in mesencephalic dopaminergic neurons shows that low ( $0.25-0.5 \mu$ M) concentrations of 1-methyl-4-phenylpyridinium (MPP+), the active metabolite of MPTP, induce neuritic tree collapse without loss of cell viability (Gómez et al. 2010). This collapse was effectively prevented by decreasing iron supply or by the addition of antioxidants. Thus, it seems plausible that increased intracellular iron and ROS are involved in the early steps of dopaminergic neuron dysfunction, prior to cell death. Later, a vicious cycle of further iron accumulation, complex I dysfunction, and ROS increase may result in uncontrolled oxidative damage and cell death.

Iron toxicity is not restricted to dopaminergic neurons. Neurotoxic concentrations of NMDA induce the NO-Dexras1-PAP7 signaling cascade in glutamatergic PC12 cells. Upon activation, PAP7 binds to intracellular DMT1 and relocates it to the plasma membrane, increasing intracellular iron and the production of hydroxyl radicals. Thus, the DMT1-iron-uptake-hydroxyl radical signaling pathway appears to mediate NMDA neurotoxicity (Cheah et al. 2006).

## 3.2 Glutathione (GSH) Metabolism in PD

The tripeptide GSH ( $\gamma$ -L-glutamyl-L-cysteinylglycine) is the most abundant and the main antioxidant agent in the central nervous system, where it reaches mM concentrations in the cytoplasm (Meister and Anderson 1983; Dringen et al. 2000). In its redox cycling, glutathione is present either in its reduced (GSH) form or its oxidized disulfide (GSSG) form, the ratio GSH:GSSG being faithful reflection of the redox state of the cell (Schafer and Buettner 2001).

Early postmortem studies revealed decreased levels of GSH in degenerating SN of PD patients (Perry et al. 1982; Sofic et al. 1988; Sian et al. 1994), the observation implicating that GSH depletion may play a major role in the neurodegenerative process. The question arises whether GSH depletion is an early event during the progression of the disease or a reflection of increased oxidative stress resulting, for example, from mitochondrial complex I inhibition and/or iron accumulation.

Chronic submaximal inhibition of GSH synthesis in N27 dopaminergic cells results in about 50 % inhibition of mitochondrial electron transport chain complex I without ensuing cell death (Chinta and Andersen 2006). Thus, increased oxidative stress generated by complex I inhibition should result in decreased GSH levels and further inhibition of complex I. Conversely, a decrease in GSH levels, provoked by unknown causes, could result in inhibition of complex I activity.

Iron induces the consumption of GSH. After exposure to increasing concentrations of iron, SH-SY5Y dopaminergic cells undergo sustained iron accumulation and a biphasic change in intracellular GSH levels, increasing at low  $(1-5 \mu M)$  iron concentrations and decreasing thereafter. Indeed, cell exposure to high iron concentrations (20–80  $\mu$ M) markedly decreases the GSH:GSSG molar ratio and the GSH half-cell reduction potential, with the associated loss of cell viability (Núñez et al. 2004).

It is therefore possible that a decrease in GSH levels is a consequence of the increased oxidative load produced by the increased intracellular iron. Nevertheless, increased iron and decreased GSH may be intertwined in a positive feedback loop, since in dopaminergic neurons the pharmacological reduction of GSH levels results in increased levels of TfR and an increased labile iron pool (Kaur et al. 2009). Thus, the question remains as to which of the three processes initiates the oxidative spiral, but a reasonable assumption is that if one of them develops, the others will follow.

## 4 Mitochondrial Dysfunction in Parkinson's Disease

Mitochondrial dysfunction is another event involved in Parkinson pathogenesis. Evidence of mitochondrial dysfunction in PD began in the 1980s, when, after an intravenous injection of illicit drugs, four people developed a marked Parkinsonism. Further analysis of the substances injected revealed the existence of MPTP, a potent inhibitor of complex I of the mitochondrial respiratory chain (Langston et al. 1983). Since its discovery, mice and monkeys intoxicated with MPTP have been used as experimental models of PD.

MPTP crosses the blood-brain barrier and it is oxidized by astrocytes to the active metabolite MPP+. Subsequently, MPP+ is selectively taken up by dopaminergic neurons through the dopamine transporter DAT (Gainetdinov et al. 1997). Inside neurons MPP+ inhibits mitochondrial complex I (Mizuno et al. 1987). Among the effects of this inhibition are decreased levels of ATP (Scotcher et al. 1990) and increased production of free radicals and oxidative stress (Wong et al. 1999).

Decreased activity of mitochondrial complex I, found in postmortem tissue of PD patients (Schapira et al. 1990; Tretter et al. 2004; Banerjee et al. 2009; Hattingen et al. 2009), is probably a founding event in neuronal death. The association between complex I inhibition and PD is further supported by the observation that rats intoxicated with rotenone, a selective inhibitor of complex I, develop a syndrome similar to PD, characterized by neuronal degeneration and the formation of inclusion bodies rich in alpha-synuclein (Betarbet et al. 2000). Likewise, inhibition of glutaredoxin 2, an enzyme involved in mitochondrial Fe–S synthesis, produces an alteration in iron metabolism in a model of Parkinson's disease (Lee et al. 2009). Additionally, the fact that mutations in mitochondrial response to oxidative stress (Valente et al. 2004), and DJ-1, an antioxidant protein that relocalizes to mitochondria under conditions of oxidative stress (Blackinton et al. 2005), result in a genetic form of PD lends further support for an important role of mitochondria in PD neurodegeneration.

ROS seem to have a negative effect on complex I activity. Experiments with isolated synaptosomal mitochondria revealed that low concentrations of  $H_2O_2$  decrease complex I activity by 10 %. This relatively minor effect of  $H_2O_2$  was additive to partial inhibition of complex I induced by low concentrations (5 nM–1  $\mu$ M) of rotenone (Chinopoulos and Adam-Vizi 2001). Similarly, submitochondrial particles exposed to  $O_2^{\bullet}$ ,  $H_2O_2$ , or  $\bullet$ OH present decreased activity of NADH dehydrogenase, a marker of complex I activity (Zhang et al. 1990). Thus, an initial inhibition of complex I could generate a positive loop between ROS generation and further complex I inhibition.

## 5 Fe–S Cluster Synthesis and Its Relevance to PD

## 5.1 Fe–S Cluster Biogenesis

Iron–sulfur (Fe–S) clusters are small inorganic cofactors formed by tetrahedral coordination of iron atoms with sulfur groups. The Fe–S clusters most commonly found in eukaryotes are 2Fe–2S and 4Fe–4S. Fe–S clusters are cofactors for proteins that are involved in many cellular processes, including electron transport, enzymatic catalysis and regulation, and DNA synthesis (Lill and Muhlenhoff 2008). The proteins that contain Fe–S clusters in eukaryotes are present in mitochondria, endoplasmic reticulum, cytoplasm, and nucleus (Lill et al. 2006; Sheftel et al. 2010).

The mitochondrion plays a central role in the generation and biology of Fe–S clusters since it holds the assembly machinery responsible for their synthesis (Lill et al. 2006). Fe–S cluster synthesis also occurs in the cytoplasm, albeit at a minor scale (Ye and Rouault 2010). In eukaryotes, the Fe–S cluster assembly machinery of mitochondria comprises the cysteine desulfurase, Nfs1, which provides sulfur and Isu1, a protein that serves as a molecular scaffold for the assembly of the Fe–S cluster. Synthesis of the transiently bound Fe–S cluster on the Isu scaffold proteins requires reduced (ferrous) iron and the input of electrons, presumably to generate sulfide (S<sup>2–</sup>) from cysteine (Ye and Rouault 2010). Finally, synthesized Fe–S clusters either are transferred to mitochondrial apoproteins or are exported into the cytoplasm by the Fe–S cluster export machinery. This machinery involves the ABCB7 transporter, the antioxidant GSH, and the sulfhydryl oxidase Erv1, which is also involved in protein import (Lill et al. 2006; Sheftel et al. 2010). Iron is transported inside the mitochondria by the transporter mitoferrin (Shaw et al. 2006).

#### 5.2 Biological Functions of Iron–Sulfur Clusters

Mitochondria contain numerous Fe–S proteins essential for its function. Complex I has eight Fe–S clusters, succinate dehydrogenase (complex II) has three Fe–S clusters, and ubiquinone cytochrome c oxidoreductase (complex III) has one 2Fe–2S cluster. Additionally, mitochondrial aconitase contains a 4Fe–4S cluster in its catalysis center (Beinert et al. 1983). Proteins involved in purine metabolism in the

cytoplasm like xanthine oxidase and phosphoribosylpyrophosphate amidotransferase (GPAT) contain two 2Fe–2S and one 4Fe–4S cluster, respectively (Unciuleac et al. 2004; Martelli et al. 2007). As mentioned above, disassembly of the 4Fe–4S cluster in IRP1 alters the active site accessibility and determines IRP1 binding to target mRNAs (Wallander et al. 2006; Cairo and Recalcati 2007). In the nucleus, proteins involved in DNA repair and replication contain Fe–S clusters (Sheftel et al. 2010). For a compendium of Fe–S clusters, see the Prosthetic Groups and Metal Ions in Protein Active Sites (PROMISE) http://metallo.scripps.edu/promise/MAIN.html.

## 5.3 IRP1: A Link Between Fe–S Clusters and Iron Homeostasis

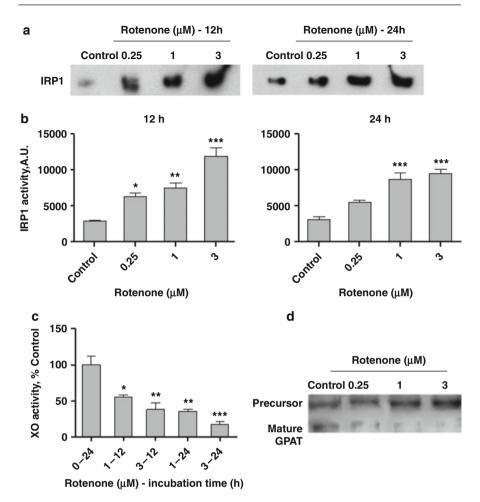
Recent data indicate that inhibition of complex I by rotenone results in decreased synthesis of Fe–S clusters, as shown by the decreased activity of the Fe–S cluster-containing enzymes cytoplasmic aconitase, mitochondrial aconitase, xanthine oxidase (XO), and GPAT, as well as the activation of cytoplasmic IRP1 (Mena et al. 2011) (Fig. 3).

It is considered that decreased activity of complex I results, via decreased Fe–S cluster synthesis and the consequent activation of IRP1, in a false "low iron" signal that activates the iron-uptake system. In consequence, diminished Fe–S cluster synthesis could play a fundamental role in the accumulation of iron observed in Parkinson's disease. Future research is needed to evaluate its participation in neurodegenerative diseases in which iron accumulation is observed.

## 6 Aminochrome: An Inhibitor of Complex I of the Mitochondrial Electron Transport Chain

The causes for complex I dysfunction in sporadic PD are a matter of active research. Exogenous toxins can be of relevance, given the existing correlation between PD and exposure to toxins such as MPTP, rotenone, paraquat, fenazaquin, annonacin, and Guadeloupe PSP-PDC (reviewed in Schapira (2010)). Endogenous toxins could also cause mitochondrial dysfunction, as is the case for nitric oxide (Clementi et al. 1998; Broom et al. 2011), 4-hydroxynonenal (Humphries et al. 1998; Castellani et al. 2002), and advanced glycosylated end products (Munch et al. 2000; Remor et al. 2011).

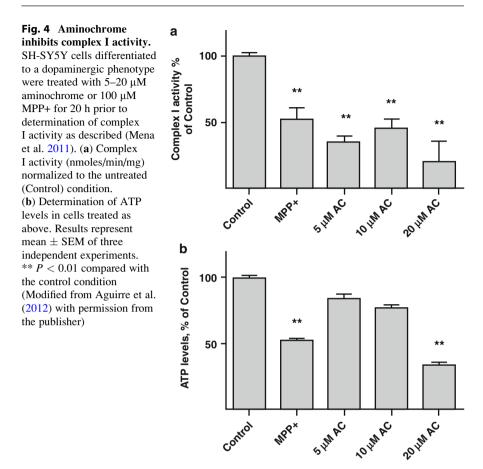
An endogenous source of free radicals proper to dopaminergic neurons derives from the nonenzymatic oxidation of dopaminoquinone to dopamine o-quinone that at physiological pH spontaneously cycles to o-quinone *aminochrome* (Arriagada et al. 2004; Zoccarato et al. 2005). A one-electron gain by aminochrome results in the highly reactive leukoaminochrome o-semiquinone radical that reacts with oxygen to generate  $O_2^{\bullet}$  (Segura-Aguilar et al. 1998). Moreover, aminochrome seems to mediate the transfer of electrons from respiration-active complex I into oxygen, thus increasing the production of  $O_2^{\bullet-}$  and  $H_2O_2$  (Zoccarato et al. 2005). The increase in reactive oxygen species has direct



**Fig. 3** Complex I inhibition affects IRP activity, XO activity, and Fe–S cluster content of GPAT. (a) EMSA for IRP activity. Cells were treated for 24 h with 0.25, 1, or 3  $\mu$ M rotenone. Subsequently, the cytosolic extracts were probed with an IRE sequence tagged with biotin. (b) Densitometric quantification of data in A. Data represent mean  $\pm$  SEM; n = 3. (c) XO activity of cells treated for 12 or 24 h with 0.25, 1, or 3  $\mu$ M rotenone. Data represent mean  $\pm$  SEM; n = 3. (d) Cells were treated for 24 h with 0.25, 1, or 3  $\mu$ M rotenone. Subsequently, the holo and apo forms of GPAT were detected by Western blot. \* *P* < 0.05; \*\* *P* < 0.01; \*\*\* *P* < 0.001 compared with the control condition (Modified from Mena et al. (2011) with permission from the publisher)

consequences on the mitochondrial electron transport chain since both  $H_2O_2$ and the 'OH inhibit complex I activity and decrease ATP production (Zhang et al. 1990; Bautista et al. 2000; Sims et al. 2000; Chinopoulos and Adam-Vizi 2001).

Recent work from our laboratory has established that aminochrome inhibits complex I and ATP production in SH-SY5Y neuroblastoma cells differentiated into a dopaminergic phenotype (Fig. 4) (Aguirre et al. 2012).



The inhibition is apparently direct on the mitochondria and not secondary to other process since it was replicated in isolated mitochondria. Additionally, overnight treatment with low micromolar concentrations of aminochrome induced increased expression of the iron import transporter DMT1 and a decrease in the export transporter FPN1. Both changes resulted in increased iron uptake. These results strongly suggest that aminochrome promotes complex I inhibition and increased iron accumulation that ends up in cell death. Therefore, aminochrome should be considered an endogenous toxin that, if uncontrolled, can induce deregulation of iron homeostasis.

## 7 Inflammation, Hepcidin, and PD

In addition to iron accumulation, another event strongly associated with neuronal death in PD and other neurodegenerative disorders is the presence of inflammatory processes characterized by the occurrence of reactive microglia and massive production of pro-inflammatory cytokines. Both phenomena have been studied as

independent events leading to the progression of the disease, but, as discussed above, the expression of iron homeostasis proteins such as ferritin, DMT1, and Dcytb is regulated by inflammatory cytokines. Moreover, the recent identification of hepcidin in the central nervous system reinforces the idea that in PD a chronic inflammation state and deregulation of iron homeostasis are intimately related.

#### 7.1 Hepcidin, the Master Regulator of Iron Homeostasis

Hepcidin is a 25-amino acids cationic peptide of secreted into bloodstream by the liver. Initially, hepcidin was described as a peptide with antimicrobial activity (Krause et al. 2000); however, further studies revealed that it also acts as a major regulator of circulating iron levels (Nicolas et al. 2001; Pigeon et al. 2001). Two processes contribute to blood iron levels: the recycling of senescent red blood cells (RBC) and intestinal iron absorption. The recycling by spleen macrophages of heme iron from senescent RBC is the main contributor of iron to the circulation, providing about 95 % of daily turnover. The recycling of RBC iron comprises the phagocytosis of senescent RBC, the release of iron contained in the heme moiety of hemoglobin by heme oxygenase-1, and the subsequent iron release into the blood mediated by FPN1 (De Domenico et al. 2008; Kovtunovych et al. 2010).

The physiological function of hepcidin is to reduce the levels of circulating iron by downregulation of the iron exo-transporter FPN1 in macrophages. The binding of hepcidin to FPN1 present in the plasma membrane of splenic macrophages induces the endocytosis of the complex and the subsequent lysosomal degradation of FPN1 (Nemeth et al. 2004). The decreased levels of FPN1 lead to the accumulation of iron in macrophages and the decrease of circulating iron.

Hepcidin synthesis is regulated by multiple stimuli that have an effect in the regulation of circulating iron levels. Erythropoietin, a hormone that stimulates red blood cell production, and hypoxia downregulate the synthesis of hepcidin, increasing circulating levels of iron as required for the production of new RBCs (Peyssonnaux et al. 2007; Pinto et al. 2008). On the contrary, increased plasma iron levels and inflammatory cytokines stimulate hepcidin synthesis, which results in reduced levels of plasma iron and its accumulation in cells (Lee et al. 2005; Wrighting and Andrews 2006; Gnana-Prakasam et al. 2008; Frazier et al. 2011).

#### 7.2 Hepcidin Expression in the CNS

Hepcidin shows a wide distribution in the CNS, most notably in the midbrain, with a clear presence in the superior colliculus, the geniculate nucleus, some fiber bundles of the SN pars reticulata, the SN pars compacta, and the striatum (Zechel et al. 2006) (Wang et al. 2010b). Hepcidin is expressed mainly in glial cells, as well as in endothelial cells of the choroid plexus (Zechel et al. 2006; Marques et al. 2009). Hepcidin expression changes with age: increased mRNA levels of hepcidin in cortex, striatum, and hippocampus have been observed with aging (Wang et al. 2010b).

As stated above, hepcidin synthesis is induced by inflammatory stimuli. Bacterial lipopolysaccharide (LPS) induces liver hepcidin expression. LPS also increases hepcidin expression in the brain. After an intraventricular injection of LPS, a transient transcription of the gene for hepcidin ensues in the choroid plexus, which correlates with increased levels of pro-hepcidin in the cerebrospinal fluid (Marques et al. 2009). The highest hepcidin expression was observed at 3 h, returning to baseline levels 24 h after the injection. Interestingly, LPS treatment induced a tenfold increase in hepcidin expression in the SN (Wang et al. 2008), which correlates with a marked increase in iron levels observed in this region in PD. Recently, it was reported that in ischemic brain there is an increase in hepcidin expression in cerebral cortex, hippocampus, and corpus striatum, in parallel with increased levels of free iron and ferritin (Ding et al. 2011). These data suggest that hepcidin could be an important contributor to iron accumulation in the brain.

#### 7.3 FPN1–Hepcidin Interactions in the CNS

As described above, the iron transporter FPN1 is a receptor for hepcidin. An interesting fact is that the spatial distribution of FPN1 and hepcidin is similar. Although the effects of hepcidin on FPN1 levels differ according to cell type (Chaston et al. 2008), the injection of hepcidin in mice lateral cerebral ventricle causes a decrease in the levels of FPN1 in the cerebral cortex, hippocampus, and striatum (Wang et al. 2010b). These results suggest that in the brain, hepcidin generates the same response as that observed in macrophages, that is, iron retention inside the cells. This conclusion is strengthened by the fact that high doses of hepcidin produce an increase in the iron-storage protein ferritin, thus indicating increased cellular iron concentration in these brain areas. Unexpected for a high cell iron situation, treatment with hepcidin induces the decrease of both FPN1 protein and mRNA and an increase in total DMT1 in the hippocampus and cortex of rats (Li et al. 2011), a situation that should drive further iron accumulation. Hippocampal neurons in culture treated with hepcidin also show a decrease in the expression of FPN1, which is reflected in a reduction of the iron released from these cells (Wang et al. 2010b). In astrocytes, one study recently reported that treatment with hepcidin significantly reduces iron uptake (both Tf-Fe and NTBI) and iron release, accompanied by decreased expressions of TfR1, DMT1, and FPN1 (Du et al. 2011).

#### 7.4 Hepcidin: A Nexus Between Inflammation and Iron Accumulation in PD

Many cases of PD are accompanied by general inflammation of the brain, with a dramatic proliferation of reactive amoeboid macrophages and microglia in the SN (McGeer et al. 1988). In the striatum, macrophage proliferation is accompanied by high expression of pro-inflammatory cytokines such as TNF- $\alpha$ , IL-1 $\beta$ , IFN- $\gamma$ , and

IL-6 by glial cells (Mogi et al. 1994; Hirsch et al. 1998; Muller et al. 1998). The increase of many of these cytokines has also been observed in cerebrospinal fluid and the basal ganglia of patients with PD (Nagatsu 2002). The aforementioned characteristics are also observed in several animal models of PD, such as those injected with 6-hydroxydopamine, MPTP, rotenone, or LPS (Herrera et al. 2000; Barnum and Tansey 2010).

The induction of hepcidin synthesis by cytokines suggests that in PD high brain hepcidin levels may mediate iron accumulation. For example, an increase in hepcidin secretion in PD triggered by inflammatory stimuli could be responsible for the decrease of FPN1 observed in this neurodegenerative condition and the subsequent accumulation of iron in neurons and glia (Lv et al. 2011).

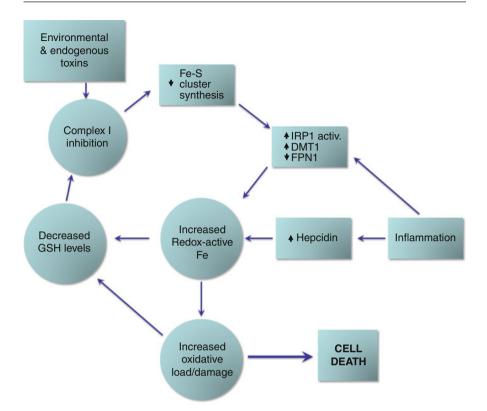
#### 7.5 Hepcidin-Independent Relationships Between Iron Accumulation and the Inflammatory Response

Inflammatory cytokines modulate the activity of IRP1 and IRP2 through nitric oxide-dependent and nitric oxide-independent mechanisms, thus influencing iron homeostasis (Mulero and Brock 1999). Stimulation of macrophages with IFN- $\gamma$ , TNF- $\alpha$ , and LPS produces an increase in IRE-binding activity of IRP1 and IRP2, reducing FPN1 and increasing DMT1 mRNA expression. Interestingly, pretreatment with the anti-inflammatory cytokine IL-10 reverts this response (Wardrop and Richardson 2000; Yang et al. 2002; Ludwiczek et al. 2003; Wang et al. 2005). In addition, DMT1 expression is transcriptionally enhanced by the transcription factor NF $\kappa$ B (Paradkar and Roth 2006a), whose activation is downstream of many cytokine receptors such as TNF receptor (TNFR) and IL-1 receptor (IL-1R). Activation of NFkB by inflammatory stimuli may play a significant role in iron accumulation by dopaminergic neurons of the SN, which express high levels of TNFR (Boka et al. 1994). Interestingly, an increase in the nuclear immunoreactivity of NF $\kappa$ B has been observed in PD brains or in animal models for this disease (Hunot et al. 1997), so it is possible that activation of NF $\kappa$ B contributes to iron accumulation in PD.

Finally, the iron transporter Zip14 could be important in the regulation of brain iron homeostasis. Highly expressed in the brain (Girijashanker et al. 2008) and upregulated by inflammatory mediators (Liuzzi et al. 2005; Lichten et al. 2009), Zip 14 could additionally contribute to iron accumulation in dopaminergic neurons of PD brains.

#### 8 A Positive Feedback Loop in the Death of Neurons

It is proposed that inhibition of mitochondrial complex I by endogenous and/or exogenous toxins and inflammatory processes produced by trauma or other causes result in a vicious cycle of increased oxidative stress, increased iron accumulation, and decreased GSH content (Fig. 5).



**Fig. 5** A positive feedback loop resulting in uncontrolled oxidative load. Complex I inhibition by endogenous or exogenous toxins results in decreased Fe–S cluster synthesis (see text). Decreased Fe–S cluster synthesis leads to the activation of IRP1 that needs a 4Fe–4S cluster to acquire its inactive state. Increased IRP1 activity results in increased DMT1 and decreased FPN1 synthesis, which produce increased iron accumulation. Increased iron induces increased oxidative stress and increased GSH consumption. Decreased GSH produces further complex I inhibition (From Núñez et al. (2012) with permission from the publisher)

In this scheme, neuronal death linked to complex I dysfunction is brought about by a positive feedback loop in which complex I inhibition results in decreased Fe–S cluster synthesis, IRP1 activation, increased DMT1 and TfR expression, and iron accumulation. Complex I dysfunction and increased cellular iron result in decreased GSH levels. Both increased oxidative stress and low GSH levels further inhibit complex I activity. Another input to this cycle is contributed by inflammatory cytokines that induce hepcidin synthesis, which by inducing FPN1 degradation results in increased neuronal iron content. Inflammatory cytokines also transcriptionally regulate DMT1, FPN1, and Zip 14 synthesis and activate IRPs (see text). Central to this scheme is the deregulation of iron homeostasis since iron chelators effectively block cell death and prevent early events in neurodegeneration such as neuritic tree shortening (Zhu et al. 2007; Gómez et al. 2010).

#### 9 Conclusion

Diminished activity of mitochondrial complex I, iron accumulation, oxidative stress, and inflammation are pathognomonic signs of sporadic PD. It is possible that the initiation of any one of these processes will initiate and enhance the others through the generation of a positive feedback loop that will produce apoptotic neuronal death. Intervention of this positive loop should result in prolonged life of the affected neurons. Still unanswered is the question of why SN pars compacta neurons are so particular prone to this deregulation.

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# Links Between Paraquat and Parkinson's Disease

Rosa A. González-Polo, José M. Bravo-San Pedro, Rubén Gómez-Sánchez, Elisa Pizarro-Estrella, Mireia Niso-Santano, and José M. Fuentes

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#### Abstract

Parkinson's disease (PD) is currently regarded as the most common degenerative disorder of the aging brain after Alzheimer's dementia. Much progress has been made in identifying the genes involved in familial, or inherited, PD. However,

R.A. González-Polo (⊠) • J.M. Bravo-San Pedro • R. Gómez-Sánchez • E. Pizarro-Estrella • M. Niso-Santano • J.M. Fuentes

Departamento de Bioquímica y Biología Molecular y Genética, Facultad de Enfermería y TO, Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas (CIBERNED), Universidad de Extremadura, Cáceres, Spain

e-mail: rosapolo@unex.es; chemabsp@unex.es; rubengs@unex.es; epizest@unex.es; mireianiso@hotmail.com; jfuentes@unex.es

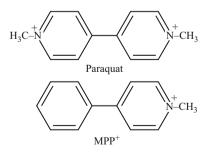
the majority of cases are sporadic (not inherited) and their origin(s) still remain largely undetermined. The environment is a key contributor to human health and disease. Epidemiological evidence suggests that environmental factors play a role in the etiology of neurodegenerative diseases. Particularly, paraquat (PQ) has been largely demonstrated to induce cell death in a variety of cell types and tissues associated with PD. The study of PQ-induced neurotoxicity has provided valuable insight into the mechanisms regulating neuronal cell death by environmental toxicants. However, to date, the molecular mechanisms involved in neuronal cell death by PQ have not been completely identified. This article presents a comprehensive review of the published epidemiologic and toxicologic literature and critically evaluates whether a relationship exists between PQ exposure and PD.

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	List of Abbreviation	ons
	ARE	Antioxidant responsive element
	ASK1	Apoptosis signal-regulating kinase 1
	CMA	Chaperone-mediated autophagy
	DA	Dopamine
	DOPAC	3,4-Dihydroxyphenylacetic acid
	DβHB	D-β-hydroxybutyrate
	ER	Endoplasmic reticulum
	ERK	Extracellular signal-regulated kinase
	GSH	Glutathione
	HVA	Homovanillic acid
	IRE1	Inositol-requiring enzyme 1
	JNK	c-Jun NH <sub>2</sub> -terminal kinase
	LAMP-2A	Lysosomal-associated membrane protein-2A
	LD	Lethal dose
	MAO-B	Monoamine oxidase B
	Mn <sup>2+</sup> -EB-DTC	Mn <sup>2+</sup> -ethylene-bis-dithiocarbamate
	MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine
	Nrf2	Nuclear factor (erythroid-derived 2)-like 2
	PD	Parkinson's disease
	PEG 400	Polyethylene glycol 400
	РКС	Protein kinase C
	PQ	Paraquat
	REP1	Dinucleotide repeat sequence
	ROS	Reactive oxygen species
	SAPK	Stress-activated protein kinase
	SOD	Superoxide dismutase
	TH	Tyrosine hydroxylase
	Trx	Thioredoxin
	WT	Wild-type

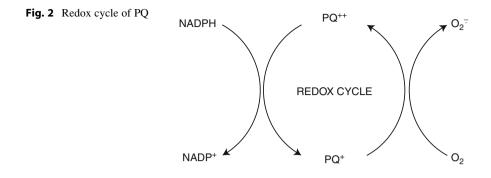
#### 1 Introduction

Paraquat (PO, methyl viologen), 1,1'-dimethyl-4,4'-bipyridinium (Fig. 1), is a widely used potent herbicide. It was first synthesized in 1882 by Weidel and Russo, as recorded by Haley in his review (Haley 1979). Its redox properties were discovered by Michaelis and Hill in 1933 (Haley 1979). At first, it was used as an indicator of oxidation reduction, since, in the absence of molecular oxygen, donating an electron to PQ ( $PQ2^+$ ) generated a monocationically stable violet or blue form, and is, thus, commonly known as methyl viologen (Dinis-Oliveira et al. 2008). But its properties as a herbicide were not discovered until 1955 and, some years later, in 1962, it was actively introduced to the global markets. PQ is registered and used in approximately 100 countries worldwide, being the second most widely used herbicide in the world after glyphosate. Despite this, today, its use is banned in the European Union (EU), unlike the import of products from outside the EU that have been treated with PQ. In its recommended rating of "pesticides by risk," the OMS (World Health Organization) considers composite PQ to be moderately toxic (Category II). The ECB (European Chemicals Bureau) classifies PO by the route of exposure, so it is considered very toxic (R26) by inhalation, toxic (R25) orally, and moderately toxic (R24) dermally. PO is included in the family of herbicides known as bipyridines. It is a contact herbicide, non-selective, systemic, and does not act on the leaves of green plants. Among its advantages, it is rapidly absorbed by the leaves of plants that are sprayed, exerting its effect very quickly, as the clay soil, which makes it biologically inactive.

There is controversy surrounding the use of PQ in agriculture because, even though its herbicidal properties are excellent, otherwise, it has clear toxicity to humans and the environment, especially proper precautions are not taken. In this sense, the structural similarity between PQ and the active metabolite of the neurotoxin called MPTP (MPP<sup>+</sup>, Fig. 1), widely accepted as a model of parkinsonism, together with some correlations observed in epidemiological studies between the use of PQ and the development of Parkinson's disease (PD) (Tanner et al. 2011), led to the postulation of the existence of a relationship between the pesticide and the origin of the disease (Costello et al. 2009; Di Monte et al. 1986; Hertzman et al. 1990;



**Fig. 1** Chemical structures of PQ and MPP<sup>+</sup>



Liou et al. 1997). Both show neurotoxic effects based on the generation of oxidative stress, but by different routes (Richardson et al. 2005). PQ is currently considered as a valid model for studying neurotoxicity based on oxidative stress, as in the case of MPP<sup>+</sup>, and investigations are ongoing concerning the relationship that may exist between application and exposure to this pesticide and the development of PD, which is a widely accepted fact in the case of MPP<sup>+</sup>, leading to an accumulation of very strong and important evidence in the case of PQ, oxidative stress, and cell death. The same active ingredient that makes PO a perfect herbicide also makes it a perfect toxic for any mammalian cell. The redox cycling of PO (Fig. 2) on biological systems has two important implications: one is the generation of reactive oxygen species (ROS) and the other is the depletion of reducing equivalents (NADH, NADPH, etc.), which are necessary for the proper functioning thereof, affecting different cellular processes, such as the synthesis of fatty acids. Like inside plant cells, within the model's neurons, PQ needs a donor of electrons in order to be reduced. The potential standard reduction ( $E^{\circ}$ ) of a compound indicates the affinity of this structure to accept electrons. PQ has an  $E^{\circ}$  of -0.45 V. The potential  $E^{\circ}$  of the redox couples NAD<sup>+</sup>/NADH and NADP<sup>+</sup>/NADPH are -0.32 V and -0.324 V, respectively, which means that PQ, under physiological conditions and with the aid of diaphorase cells, could accept electrons from either of the reducing agents. The potential  $E^{\circ}$  of MPP<sup>+</sup> is -1.18 V, which indicates that PQ has a greater ability to accept electrons than MPP<sup>+</sup> (Drechsel and Patel 2008). Among the cellular enzymes that can give electrons to PQ (PQ enzymes with diaphorase), those that can be highlighted are the following: mitochondrial complex I (NADH: ubiquinone reductase complex) (Fukushima et al. 1993), thioredoxin reductase (Gray et al. 2007), NADPH:ferredoxin oxidoreductase (Liochev et al. 1994), NADPH oxidase (Bonneh-Barkay et al. 2005), and NOS (Patel et al. 1996), as well as others. The mitochondria, therefore, becomes a major source of ROS generation within the PO-induced mechanism, which may induce PO-diaphorase activity at one or more points during the breathing cycle (Drechsel and Patel 2008). Once the PQ has been reduced, it can then be oxidized by a molecule of oxygen and generate a superoxide molecule  $(O_2^{\bullet})$ , which occurs in cell oxidative stress and will switch to different routes that trigger cell damage in different components and initiate the activation of different cellular mechanisms, such as apoptosis or autophagy. Apoptosis induced by PQ has been demonstrated to involve mainly the intrinsic mitochondrial pathway. The intrinsic pathway of apoptosis, also referred to as the mitochondrial pathway, is activated by a wide variety of cytotoxic stimuli or environmental stressors. Although the mechanisms by which these stimuli trigger apoptosis differ between them, they convey the release of pro-apoptotic proteins from the mitochondria, including cytochrome c. However, the exact mechanisms mediating cytochrome c release are still a matter of debate (Grimm and Brdiczka 2007; Ravagnan et al. 2002). In PD, cell death by apoptosis has been proposed to result from mitochondrial dysfunction, leading to an increase in oxidative stress and a decline in ATP production. PQ-induced cytotoxicity has been suggested to be mediated through the extrinsic pathway of apoptosis by activation of the Fas receptor (Vogt et al. 1998), but its relevance to the progression of PD remains unclear. On the other hand, in recent years, converging evidence suggests that the impairment of homeostasis mechanisms processing unwanted and misfolded proteins plays a central role in the pathogenesis of PD (Olanow 2007). Impairment of the autophagy-lysosomal pathway was related to the development of PD (Pan et al. 2008). The activation of autophagy was also identified within peripheral blood mononuclear cells from PD patients (Prigione et al. 2010). In this sense, PQ is able to also induce autophagy in dopaminergic cellular models (González-Polo et al. 2007b). Therefore, PQ is considered as one of the most useful models for studying neurotoxicity based on the generation of oxidative stress (such as PD), playing a key and fundamental role in the production of superoxide anions in the redox cycling of the herbicide, which can induce both apoptosis and autophagy processes.

## 2 Epidemiologic Evidence

As evidence emerges that several genes are involved in the pathogenesis of PD, the role of environmental chemicals in the etiology of this disease has become intensely debated. Many chemicals that are toxicologically or structurally similar to MPTP, an agent known to induce Parkinsonian symptoms in humans (Langston and Ballard 1983), could be a risk factor for PD. Several studies reported increased risk associated with exposure to either insecticides or herbicides (Butterfield et al. 1993; Firestone et al. 2005; Gorell et al. 2004; Semchuk et al. 1992) and organochloride, organophosphate, and carbamate insecticides (Seidler et al. 1996). The relationship between exposure duration and PD risk was investigated in several studies and showed a positive correlation with the duration of exposure to, and high doses of, herbicides and insecticides (Brown et al. 2006; Seidler et al. 1996). In these studies, PQ exposure was shown to be significantly associated with PD (Hertzman et al. 1990; Liou et al. 1997). A case–control study of environmental risk factors and PD in Taiwan reported evidence of a strong correlation between PD incidence and the level of PQ exposure (Liou et al. 1997). Another report from Dinis-Oliveira and coworkers showed that

occupational exposure to PQ was associated with parkinsonism in a study in British Columbia (Dinis-Oliveira et al. 2006).

PQ has been shown to interact synergistically with other pesticides and insecticides. It has been reported that exposure to a combination of maneb (a dithiocarbamate fungicide) and PQ increases PD risk, particularly in younger subjects and/or when exposure occurs at younger ages (Costello et al. 2009; Thiruchelvam et al. 2000). Combined exposure of rotenone and PQ has been linked experimentally to pathophysiological mechanisms implicated in human PD (Tanner et al. 2011). Another study in Central Valley, California, that combined ambient exposure to ziram and PQ as well as combined ambient exposure to maneb and PQ at both workplaces and places of residence showed substantially increased risk of PD. Those exposed to ziram, maneb, and PQ together experienced the greatest increase in PD risk (Wang et al. 2011).

Different groups of pesticides have also shown positive associations with PD. Case reports and post-mortem studies have described PD in persons exposed to organophosphate insecticides; organochlorine insecticides, including dieldrin; herbicides, including glyphosate and diquat; fungicides, including maneb and other dithiocarbamates (Corrigan et al. 1998; Firestone et al. 2005; Fleming et al. 1994; Kamel and Hoppin 2004; Wechsler et al. 1991). However, other studies have not found a significant association, although PD risk was still elevated (Firestone et al. 2005; Hertzman et al. 1990; Kamel and Hoppin 2004; Li et al. 2005a; Wirdefeldt et al. 2011). In addition, potential confounding factors include age, well-water consumption, farming, rural living, head trauma, smoking and caffeine intake, and infectious diseases such as nocardiosis, as all of these have been suggested to affect the incidence of PD (Berry et al. 2010). The Agricultural Health Study carried out a cohort investigation of over 79,500 agricultural workers from North Carolina and Iowa. The researchers considered various exposures, including acute poisoning episodes and chronic exposures, and took account of several confounding factors and the different farming practices. They found that the prevalence of neurological symptoms was associated with cumulative life-time exposure to pesticides, notably, organophosphates and organochlorines, and that this prevalence was mainly associated with chronic moderate exposure (Berry et al. 2010; Kamel et al. 2007).

### 3 Relation Between Environmental and Genetic Factors in Parkinson's Disease: The Paraquat and PARK Genes Connection

The etiology of PD is unknown, but it likely has a multifactorial origin, involving both genetic and environmental factors, such as exposure to PQ. The interaction of both factors is possibly responsible for the selective death of dopaminergic neurons observed in this disease. Outside of the studies that have identified human mutations as a basis for disease, the high number of individuals with sporadic PD have an unknown etiology. Although this interaction is not clear, there exist many studies with results that indicate the direct interaction of PQ with PARK genes (Table 1).

DADK 1/4		Protein name	Inheritance	Function
PARK 1/4	4q21.3-q22	α-synuclein	AD	LB component
PARK 2	6q25-27	Parkin	AR	E3 ubiquitin ligase
PARK 3	2p13	i?	AD	i?
PARK 5	4p14	UCHL-1	AD	Ubiquitin C-terminal hydrolase
PARK 6	1p35-36	PINK1	AR	Mitochondrial kinase
PARK 7	1p36	DJ-1	AR	Antioxidant protein
PARK 8	12q12	LRRK2	AD	Kinase, GTPase
PARK 9	1p36	ATP13A2	AR	ATPase, cationic transport
PARK 10	1p32	¿?	AD	ί?
PARK 11	2q36-q37	GIGYF2	AD	Receptor tyrosine kinase signaling
PARK 12	Xq21-q25	¿?	X-linked	ί?
PARK 13	2p13	HtrA2/Omi	AD	Serine protease
PARK 14	22q13.1	PLA2G6	AR	Phospholipase A2
PARK 15	22q11.2	FBXO7	AR	E3 ubiquitin ligase
PARK 16	1q32	RAB7L1	<i>;</i> ?	i?
PARK 17	4p	VPS35 (2011)	ζ?	i?
PARK 18	6р	EIF4G1 (2011)	ζ?	i?

Table 1 Summary of genetic ("PARK") loci associated with a monogenetic form of PD

¿? - unknown

One of the most important genes related to PD is PARK 1/4, which codifies to  $\alpha$ -synuclein and plays an important role in the LB formation, and could establish interactions with toxicants. In this sense, in 2001, Vladimir N. Uversky and colleagues (2001) showed how the presence of PQ produced the fibril formation of recombinant  $\alpha$ -synuclein in vitro, being the first approach that linked pesticides with PD. Interestingly, this fact is accelerated in a dose-dependent manner and the exposure to PQ increased the  $\alpha$ -synuclein levels in the ventral mesencephalon and frontal cortex of mice (Manning-Bog et al. 2002). Moreover, that laboratory provided evidence that the  $\alpha$ -synuclein overexpression protects front dopaminergic cell PQ-dependent degeneration in transgenic mice, for both human WT  $\alpha$ -synuclein and human Ala53Thr mutant  $\alpha$ -synuclein (Manning-Bog et al. 2003). Furthermore, the PQ exposure leads to the nuclear localization of  $\alpha$ -synuclein in mice midbrain, and it co-localizes with histone H3, and this event could be a mechanism to decrease the pool of free histones (Goers et al. 2003). Under macromolecular crowding conditions, the presence of PQ accelerates the  $\alpha$ -synuclein fibrillation compared with only the addition of polymers, such as PEG 400 (Munishkina et al. 2004). In an in vivo study, Thiruchelvam and colleagues (Thiruchelvam et al. 2004) treated male mice over a period of 7 weeks with combined PQ and Mn<sup>2+</sup>-ethylene-bis-dithiocarbamate (Mn<sup>2+</sup>-EB-DTC), observing that the mice which expressed human doubly-mutated  $\alpha$ -synuclein (A53T and A30P) presented a significant reduction in locomotor activity, dopamine (DA) levels, and TH+ neurons, as well as small decrease in 3.4-dihydroxyphenylacetic acid (DOPAC) and homovanillic acid (HVA) levels, but the DA turnover was elevated relative to saline counterparts. In HEK293 cells, G209A mutant  $\alpha$ -synuclein expressing cells were partially protected from the PO oxidative stress, but less than that for WT  $\alpha$ -synuclein cells (Orth et al. 2004). On the other hand, a report based in the nematode C. elegans (Ved et al. 2005) demonstrated that WT  $\alpha$ -synuclein and A53T mutant showed no difference in terms of sensitivity to PQ compared with the nontransgenic line, and the co-treatment with PQ and probucol increased the survival levels, but not when using D- $\beta$ -hydroxybutyrate (D $\beta$ HB), indicating that the free radical generation contributes to PQ toxicity, and, in this model, it is independent of the complex I inhibition. In astrocytes, the expression of PEP-1-SOD fusion protein with Tat- $\alpha$ synuclein increased the cell viability compared with cells pretreated with mutant proteins (A30P and A53T) after the PQ exposure, and the neuroprotective effect was synergistically higher than the expression of Tat- $\alpha$ -synuclein when measuring the SOD activity assay. Also, in that study, when PEP-1-SOD was cotransduced with WT Tat- $\alpha$ -synuclein, the number of TH<sup>+</sup> neurons and the Hsp70 protein levels were enhanced compared with PO-treated cells, but not when they overexpressed A53T mutant Tat- $\alpha$ -synuclein (Choi et al. 2006). A study by Fernagut et al. (2007), using mice overexpressing human WT  $\alpha$ -synuclein under the Thy-1 promoter (Thy1-aSYN), demonstrated how they had several impaired postural adjustments (such as errors per step, time to orient downward, etc.), which were unchanged after the PO treatment. However, this herbicide induced a significant increase of the proteinase K-resistant  $\alpha$ -synuclein inclusions in the substantia nigra of Thy1- $\alpha$ SYN mice, and their diameter were higher relative to the saline treatment. This fact could be due to an increased level of the protein or to the  $\alpha$ -synuclein nitration, allowing its resistance to proteolysis (Uversky et al. 2005). Studies in human neuroblastoma SH-SY5Y cells (Chau et al. 2009) have shown that PQ produced similar cell death and free radical generation in control and WT  $\alpha$ -synuclein cells; however, this rate is higher in cells overexpressing A53T mutant  $\alpha$ -synuclein. Also, the caspase-3 activity was highest in the mutant cells, in line with the mitochondrial membrane potential reduction, indicating apoptosis mediated by PQ. Otherwise, the co-incubation with cabergoline (dopamine agonist) decreased this pattern of cell death, having better protection in A53T mutant cells. Another neuroprotector was used to prevent PQ toxicity (Chau et al. 2010), rasagiline (irreversible MAO-B oxidase inhibitor), demonstrating that it ameliorated the PQ cell death in SH-SY5Y cells overexpressing or not WT or A53T mutant  $\alpha$ -synuclein, as well as decreasing the PQ-induced caspase-3 activation and free radical production. In the same way, the A53T mutant mice suffered a TH+ neuronal loss in a synergic manner with PQ, combined or not with iron, and this event is age-dependent, just as the increase of the protein oxidation levels. However, the dopaminergic neuronal death and the nitrated  $\alpha$ -synuclein can be partly rescued using the antioxidant EUK-189 (Peng et al. 2010). A recent epidemiologic report showed an approach between  $\alpha$ -synuclein and environmental factors, like PO, because of the low risk of PD development in subjects who were either homozygous or heterozygous for the α-synuclein REP1 259 genotype, and subjects who were homozygous or heterozygous for the REP1 263 genotype have a higher risk, but in PQ exposure areas, the protective effect of the shorter repeat REP1 could be lost, and the longer REP1 genotype may enhance this susceptibility and promote a higher risk in younger populations (Gatto et al. 2010). An important item could be if there is a mechanism to maintain the  $\alpha$ -synuclein levels in the cell, and some studies (Mak et al. 2010) have shown how  $\alpha$ -synuclein is associated with lysosomes in the midbrain of PQ-treated mice to uptake and internalize this protein for later clearance in this organelle via chaperone-mediated autophagy (CMA), because the PQ exposure increased the hsc70 and LAMP-2A levels and their interaction with  $\alpha$ -synuclein. To establish a relationship between CMA and  $\alpha$ -synuclein, the investigators used transgenic mice overexpressing mouse  $\alpha$ -synuclein, observing higher levels of LAMP-2A and hsc70. A new report showed that lithium protects against neuronal cell death and prevents nitrated/oxidized  $\alpha$ -synuclein accumulation in human A53T mutant-expressing mice chronically exposed to PQ/maneb in several brain regions (Kim et al. 2011).

Another gene involved in PD is *parkin (PARK2)*, and several studies have demonstrated that the environmental stress also modulates the corresponding protein. In this way, a report by Pesah and colleagues showed how *parkin* mutant flies have enhanced sensitivity to PQ toxicity (Pesah et al. 2004), claiming that parkin could have a protective role. Also, PQ changes the parkin solubility and promotes the formation of intracellular parkin inclusions in SH-SY5Y cells expressing FLAG-parkin (Wang et al. 2005). In a previously cited study (Ved et al. 2005), in *C. elegans*, the *KO8E3.7/Parkin KO* did not show a higher sensitivity to the PQ treatment, and the same pattern was demonstrated in the co-treatment with probucol and D $\beta$ HB as in the previous paragraph. In *parkin* mutant human fibroblasts, the treatment with PQ induced significant loss of mitochondrial membrane potential, slight increase in protein oxidation, and reduction of mitochondrial branching (Grünewald et al. 2010).

Following the above, *PINK1* (*PARK6*) is a gene that is strongly related to PD. In this case, in *PINK1* silenced cells, PQ generated an important reduction in cell viability and a higher rate of apoptosis cell death (Gegg et al. 2009). In primary fibroblasts cultures, the mitochondrial membrane potential was unaffected in control and mutant cells after PQ exposure, but the glutathione (GSH) levels were increased in missense culture to a greater extent (Grünewald et al. 2009). A report realized in *C. elegans* (Sämann et al. 2009) suggested a functional linkage between the *lrrk-1* and *pink-1* homologs of human *LRRK2* (*PARK8*) and *PINK1* genes, showing how the loss of *pink-1* function produced a higher sensitivity to PQ, whereas the genetic deletion of *lrrk-1* could compensate for both of these deficiencies.

Finally, DJ-1 (CAP1/RS/PARK7) is a protein related to early-onset PD, for which patients respond to treatment with dopamine (Bonifati et al. 2003). Although the function of DJ-1 is unclear, several studies suggest their possible involvement in the cellular response against oxidative stress, because DJ-1 has the capacity to eliminate ROS by auto-oxidation (Bonifati et al. 2003; Kinumi et al. 2004; Taira et al. 2004). It has been observed that, if DJ-1 shows mutations in C106, it is unable

to protect the cell under conditions of oxidative stress caused by MPP<sup>+</sup> (an inhibitor of mitochondrial complex I) (Canet-Avilés et al. 2004). These findings suggest that mutations in the DJ-1 protein can cause mitochondrial susceptibility against toxins such as MPP<sup>+</sup>. There are many cellular and animal models available to investigate the combined effect of genetic and environmental factors involved in PD and to study the interaction between DJ-1 and the herbicide PQ. The idea that DJ-1 stimulates the autophagy response against PQ and reduces apoptotic cell death response could open the door to the development of new treatment strategies against PD (González-Polo et al. 2010).

#### 4 Cellular Models in Paraquat Neurotoxicity

In PD, cell death by apoptosis has been proposed to result from mitochondrial dysfunction, leading to an increase in oxidative stress and a decline in ATP production. Some studies have shown that PQ induces cytochrome c release (Fei et al. 2008; González-Polo et al. 2004) and caspase-9 activation, which are preceded by the induction/activation of pro-apoptotic Bax and Bak (Fei et al. 2008; Yang and Tiffany-Castiglioni 2008). The evaluation of Bcl-2 family members showed that PQ induces high levels of Bak, Bid, BNip3, and Noxa (Fei et al. 2008). Interestingly, there are many works which demonstrate that, individually, PO and maneb (and other pesticides related to PD) activate Bak, but, together, they trigger Bax-dependent cell death. Focusing on mechanisms responsible for this synergy, the study showed that maneb + PO increases the expression of three strong Bak inhibitors, Bfl-1, Bcl-xL and Mcl-1, and, also, induces Bax activators that included Bik and Bim (Fei and Ethell 2008). The induction of pro-apoptotic Bax and apoptosis in response to PQ has also been reported to be dependent on p53 (Rossi et al. 2001; Yang and Tiffany-Castiglioni 2008). PQ neurotoxicity has also been reported to require the activation of stress-activated protein kinases (SAPKs) (Chun et al. 2001; Klintworth et al. 2007; Niso-Santano et al. 2006, 2010; Ramachandiran et al. 2007; Yang et al. 2009).

Although the mitochondrial pathway of apoptosis has been largely linked to PQ-induced cell death, other signaling pathways have also been implicated in PQ toxicity. Recently, PQ has been shown to induce DNA damage and endoplasmic reticulum (ER) stress. ER stress is associated with the activation of inositol-requiring enzyme 1 (IRE1), apoptosis signal-regulating kinase 1 (ASK1), and c-Jun NH<sub>2</sub>-terminal kinase (JNK) (Niso-Santano et al. 2010; Yang et al. 2009). In this sense, PQ-induced oxidation of Trx has been reported as a possible mechanism for the activation of the ASK1/JNK signaling pathways (Niso-Santano et al. 2010; Ramachandiran et al. 2007; Yang et al. 2009). NF-E2-related factor-2 (Nrf2)-dependent regulation of antioxidant responsive element (ARE)-mediated gene expression has been shown to protect against PQ (Minelli et al. 2009). Accordingly, Nrf2-dependent regulation of Trx levels determines the sensitivity of PQ toxicity by the activation of ASK1/JNKp38 signaling (Niso-Santano et al. 2010). PQ-induced tyrosine nitration and lipid peroxidation have been

demonstrated (Schmuck et al. 2002). However, the molecular targets for these signaling events remain to be elucidated. It was recently demonstrated that oxidative stress induced by PQ generates protein aggregation of the plasma membrane  $Ca^{2+}$ -ATPase and its degradation by calpain (Zaidi et al. 2009).

Recent studies have demonstrated that low concentrations of PQ induce autophagy, which is followed by apoptosis (González-Polo et al. 2007a, b, 2009) and modulated by DJ-1. Because the inhibition of autophagy potentiated apoptosis induced by PQ, it was proposed that autophagy might be acting as a protective mechanism against cell death progression.

The fact that PQ induces the accumulation of autophagic vacuoles and increases the degradation of proteins in the cytoplasm of SH-SY5Y cells (González-Polo et al. 2007a) indicates that the increased oxidative stress can activate autophagy in the initial stages of mitochondrial dysfunction, having a protective role of PQ-induced cell death (González-Polo et al. 2007a, b). Moreover, it has been shown that PQ exposure induces an early ER (endoplasmic reticulum) stress response that is correlated with this adaptive activation of autophagy, with hallmarks such as the accumulation of autophagic vacuoles, activation of beclin-1, accumulation of LC3-II, p62 degradation, and mTOR dephosphorylation (González-Polo et al. 2007a, b; Niso-Santano et al. 2011). This response is accelerated in cells that overexpress WT of the ASK1 protein. The inhibition of autophagy caused, in this model, an exacerbation of the apoptosis induced by WT ASK1 overexpression with or without PQ. These results suggest that autophagy has an important role in the cell death/survival events produced by PQ and ASK1 that contribute to neuronal degeneration.

Therefore, increased autophagy might be a new strategy for the treatment of neurodegenerative diseases (Menzies et al. 2006). However, it is encouraging to consider enhancing autophagic capacity as a therapeutic strategy in the prevention of neurodegeneration because, as mentioned above, there is evidence showing that the abnormal regulation of autophagic pathways may lead to apoptosis and cell death (Walls et al. 2010).

Most of the studies regarding the molecular mechanisms of PQ-induced apoptosis and autophagy have been directed towards neuronal cell types (Table 2). However, it is obvious that the complexity of brain tissue organization is given by the interaction of different neurons with glial cell types. Compared to neurons, astrocytes have been demonstrated to be more resistant to PQ-induced oxidative stress by the responsive induction of a variety of antioxidant systems, including catalase and SODs (Olesen et al. 2008; Röhrdanz et al. 2001; Schmuck et al. 2002). Recent studies have demonstrated that microglial cells react to pro-oxidant conditions, suggesting that they can contribute to environmental stress-induced neurotoxicity (Vogt et al. 1998). PQ induces microglial activation, which seems to precede PD neurodegeneration (Purisai et al. 2007). Low PQ concentrations have recently been demonstrated to be toxic to neurons only in the presence of microglial cells. In this study, it was observed that, in neuron-microglia cultures exposed to PQ, microglia was a source of PQ-derived oxidative stress (Wu et al. 2005). Interestingly, NADPH oxidase from glial cells has been proposed to mediate the

Model	Apoptotic major findings	References	
Cerebellar granule cells	Caspase 3 activation	González-Polo et al 2004	
	Cytochrome c release		
	DNA fragmentation		
Rat brain neuroblasts, E18 cells	ERK1/2 activation	Niso-Santano et al. 2006	
	JNK activation		
	PKB activation		
Pheochromocytoma PC12 cells	JNK/p38 activation	Klintworth et al. 2007	
Rat primary cultured dopaminergic neurons	JNK activation	Klintworth et al. 2007	
SK-N-SH neuroblastoma cells	Bak activation	Fei et al. 2008	
	Cytochrome c release		
	Caspase 3 activation		
	Thioredoxin oxidized	Ramachandiran	
	JNK activation	et al. 2007	
	Caspase 3 activation		
Substantia nigra-derived	JNK/MAPK activation	Chun et al. 2001	
dopaminergic SN4741 cell line	Caspases 1 and 3 activation		
SH-SY5Y human neuroblastoma cells	JNK/p38 activation	Klintworth et al. 2007	
	Mitochondrial intrinsic pathway associated with p53	Yang and Tiffany- Castiglioni 2008	
	IRE1/ASK1/JNK activation	Yang et al. 2009	
	ASK1/JNK activation	Niso-Santano et al 2010	
	ASK1/p38 activation		
	Caspase 3 activation		
Autophagic major findings			
SH-SY5Y human	Accumulation of autophagic vacuoles	González-Polo et al	
neuroblastoma cells	Recruitment of an LC3-GFP fusion protein to autophagic vacuoles	2007a, b	
	Long-lived protein degradation regulated by mTOR signaling	88	
	ER stress response	Niso-Santano et al	
	Beclin-1 activation		
	p62 degradation		
	Autophagic events DJ-1 dependent	González-Polo et al 2009	

Table 2 List of studies on the induction of apoptosis and/or autophagy by PQ

generation of ROS (Kim et al. 2008; Purisai et al. 2007; Schmuck et al. 2002), which seems to be regulated by the activation of protein kinase C (PKC) and extracellular signal-regulated kinase (ERK) signaling pathways (Kim et al. 2008; Miller et al. 2007).

#### 5 Animal Models in Paraquat Neurotoxicity

Animal models are an important tool for studying different human diseases, including PD. Animals do not develop PD spontaneously, but it is possible mimic the symptoms of this disease, either by surgery or the use of chemicals to damage the tissues of the body in a way that reflects the changing disease. Animal models allow scientists prove what is causing parkinsonism symptoms and to test different treatments that cannot be tested in people. Therefore, despite these models never being able to fully mimic the human model, they are vital in order to understand the molecular and biochemical features of PD (Orth and Tabrizi 2003).

Using drugs to alter the animals is the main method used in research on Parkinson's disease. MPTP, rotenone, or PQ are used to cause parkinsonian symptoms in different animal models. The animal models commonly used for PD investigations are monkey, rat, or mice, but animals like rabbits, fruit flies, dogs, nematodes, zebrafish, pigs, or guinea pigs are also used, although to a lesser extent.

These models have highlighted the possibility that environmental chemicals used in everyday situations may be responsible for contributing to the development of PD. Behavioral effects and loss of neurones in the substantia nigra were observed and the depletion of dopamine was reported in many, but not all, of these studies.

The following is a review of the different animal models used for the studies of parkinsonian symptoms associated with exposure to PQ.

#### 5.1 Rat Animal Model

The use of a rat animal model to study parkinsonian symptoms is very common. Exposure to PQ in rats induces much damage at different levels in rats, and presents lung and renal toxicity and toxicity to the liver. At higher doses, observations include decreased body weight gain, clinical signs like dyspnea, increased respiratory sounds, swellings and sores in the genital area, hematological changes, and effects on organ weight, as well as increased mortality.

In addition, a large number of studies have been performed in rat brains to observe animal behavior after being treated with PQ. It has been shown that there are changes in behavior as well as neuropathological effects in rats which received both systemic and intrahippocampal injections of PQ dichloride (Corasaniti et al. 1992). After treatment with PQ by unilateral injection into the substantia nigra, there was loss of Nissl substance, glial reaction, and loss of neurones in the substantia nigra, and, neurochemically, there was dopamine depletion (Liou et al. 1996). The neuropathological findings in another study showed that PQ possesses marked neurotoxicity, with neuropathological lesions culminating in neuronal necrosis (Calò et al. 1990). Another study showed that PQ induces oxidative stress, neuronal loss in the substantia nigra region, and parkinsonism in adult rats (Somayajulu-Niţu et al. 2009) and lipid peroxidation in rat brains (Yumino et al. 2002). This neurodegeneration is observed after exposure to PQ, which might occur via an early inflammatory response in young adult animals (Cicchetti et al. 2005).

Therefore, in rats, it has been shown that PQ induces neuronal damage and changes in the behavior of these animals. Furthermore, this damage could be caused by oxidative stress, lipid peroxidation, and inflammatory responses after stimulation with PQ.

#### 5.2 Mouse Animal Model

Many in vivo studies in mice have shown that acute PQ administration may elicit a distinct decrease in the level of dopamine and spontaneous motor activity in mice (Kang et al. 2009; Li et al. 2005b) and there are studies that relate directly the PQ exposure in mice with a decrease in motor activity (Brooks et al. 1999). However, equivocal results have been reported by studies investigating the effect of exposure to PQ on selective nigrostriatal degeneration in rodents. There are studies which show that PQ does not induce Parkinson's disease symptoms in mice (Rojo et al. 2007), but other works show that PQ administration triggers an apoptotic cell death program through oxidative stress-mediated activation of the JNK signaling pathway, suggesting a possible mechanism for selective dopaminergic neuron loss (Peng et al. 2004). There has been a study on C57BL/6 mice exposed to PQ, maneb, or a PQ + maneb mixture, a combination that produces PD symptoms in mice (Thiruchelvam et al. 2000).

With a better understanding of the genetic alterations involved in familial PD, several transgenic mouse models have been generated that can reproduce many features of parkinsonism. Animal models generated by mutations or deletions of the relevant Parkinson's-related genes have also been used to examine the effects of PQ; for example, studies have shown that the overexpression of  $\alpha$ -synuclein protects against PQ toxicity (Manning-Bog et al. 2002, 2003).

#### 5.3 Monkey Animal Model

Monkey species, like *Saimiri sciureus* or *Macaca fascicularis*, are normally used in studies with PQ treatments. There have been studies carried out to characterize and compare the toxicity, clinical syndromes, and pathology of PQ, using an LD<sub>50</sub> in these animals of normally 50–70 mg/kg body weight. Many decades ago, there were studies indicating that the oral administration of PQ in this species induced clinical signs of intoxication, such as anorexia, adipsia, diarrhea, hyperpnea, dyspnea, and tachycardia, which depend on the duration of exposure and the concentration of PQ (Murray and Gibson 1972; Purser and Rose 1979). However, there are few studies of monkeys concerning the neuronal damage caused by PQ exposition, although recent studies on PQ administration in monkeys have demonstrated that it produces a depletion of dopamine in the striatum of these animals (O'Leary et al. 2008).

#### 5.4 Fruit Fly Animal Model

*Drosophila melanogaster* is a popular study species in science. This particular species has become an invaluable tool to scientists, who have been studying this insect for over a century. The fruit fly *D. melanogaster* has a similar number of genes to humans and, therefore, it is often used in genetic changes in order to observe the effects on the progression of disease. A large number of human genes, including those implicated in PD such as parkin, UCH-L1, PINK1, DJ-1, and LRRK2, have highly conserved homologs in *D. melanogaster* (Whitworth et al. 2006).

The exists many models of fruit flies to study genetic changes, which has shown dysfunction of DA neurons, locomotor deficits, and retinal degeneration after treatment with PQ. The  $\alpha$ -synuclein (Takahashi et al. 2003) or parkin models in flies show that PQ induces DA neuron loss, mitochondrial pathology, locomotor deficits, LB-like protein aggregates, and sensitivity to oxidative stress (Greene et al. 2005). Another study in flies indicates that DJ-1 plays a critical role in the survival of dopaminergic neurons and response to oxidative stress in PQ treatment (Menzies et al. 2005). In *D. melanogaster*, there are also studies that show damage at the level of oxidative stress which occurs with PQ exposure, along with neuronal damage that this entails (Rzezniczak et al. 2011).

#### 5.5 Nematode Animal Model

*Caenorhabditis elegans* is a nematode animal model used commonly in scientific studies. *C. elegans* models have a well-characterized genome and the adult wild-type worm comprises 1,000 cells, about a third of which are neurons, exactly eight of which are dopaminergic neurons. It also has all the dopamine transmission mechanisms of a human model (Nass et al. 2001). These dopamine neurons of *C. elegans* are sensitive to neurotoxins and pesticides like 6-OHDA (Nass et al. 2002), MPP<sup>+</sup> (Braungart et al. 2004), and rotenone or PQ (Ved et al. 2005). This has led to a search for genetic patterns among nematodes of resistance to the toxicity of PQ in genes that seem to be involved in the function of chemosensory cilia (Fujii et al. 2004, 2005) or with antioxidant genes (Yanase et al. 2002).

#### 5.6 Zebrafish Animal Model

The *Danio rerio* species is a 3–4-cm-long zebrafish used to study development and gene function as a potential model of PD. In zebrafish, dopaminergic neurons are anatomically comparable with the nigrostriatal tract in mammals. These neurons are sensitive to some of the classical PD model toxins, namely, 6-OHDA and MPTP, showing reduced brain levels of dopamine after treatment (Anichtchik et al. 2004).

A study among zebrafish with PQ showed that, in adult zebrafish, there was no induced effect after a long period of exposure to pesticide PQ, and in larval zebrafish, only a trend of reduction in diencephalic DA neurons was shown, which did not reach statistical significance (Bretaud et al. 2004).

## 5.7 Dog Animal Model

In many studies with dogs, principally the Beagle dog breed, the animals received PQ as a dietary supplement and the results indicate that levels of pulmonary capillary and pulmonary vascular resistance were increased, mean arterial pressure and cardiac output were decreased, while the total vascular resistance and pulmonary vascular resistance were increased (Sone et al. 1989). Another study showed the appearance of dyspnea, and, often, emaciation or inappetence, with some body weight loss. However, there are no conclusive studies on the neurological damage produced in dogs after exposure to PQ.

## 5.8 Rabbit Animal Model

Rabbits are another animal model used to probe the toxicity of PQ. After PQ exposition, damage by pneumonitis has been detected in the animals (Seidenfeld et al. 1985), as well as damage to the eyes, liver, kidneys, heart, and lungs (Hassan et al. 1989), but there exists no studies demonstrating damage at the neurological level.

# 5.9 Pig Animal Model

Pigs have been used on rare occasions for PQ toxicity studies, showing that high concentrations of and prolonged exposure to PQ may have deleterious effects on epidermal morphology in the absence of significant percutaneous absorption (Srikrishna et al. 1992), but there has been no study neuronal damage in these animal models.

# 6 Conclusion

Toxicologic studies have suggested that multiple genetic and environmental factors could be involved in the etiology of PD. Studies with transgenic mice suggest that the genetic background may also lead to increased vulnerability to the neurotoxic effects of pesticides such as PQ. The weight of evidence is sufficient to conclude that a generic association between PQ exposure and PD exists, but it is not sufficient to conclude that this is a causal relationship. In addition, the multifactorial etiology of PD hampers unequivocally establishment of the role of any individual contributory causal factor. Further research is needed in order to identify long-term biomarkers of exposure, improve methods for estimating pesticide exposure, and

undertake prospective cohort studies of PQ-exposed people. It is also necessary to increase the volume of studies on the molecular basis of PQ toxicity in order to establish a clear relationship between its neurotoxic effects and in vivo neuronal loss that leads to PD.

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# **Manganese Neurotoxicity**

Daiana Silva Ávila, Robson Luiz Puntel, Vanderlei Folmer, João Batista Teixeira Rocha, Ana Paula Marreilha dos Santos, and Michael Aschner

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D.S. Ávila (🖂) • R.L. Puntel • V. Folmer

Universidade Federal do Pampa, Uruguaiana, RS, Brazil e-mail: avilads1@gmail.com; robsonunipampa@gmail.com; vandfolmer@gmail.com

J.B.T. Rocha Universidade Federal de Santa Maria, RS, Brazil e-mail: jbtrocha@yahoo.com.br

A.P.M. dos Santos Department of Toxicology and Food Sciences, Faculty of Pharmacy, University of Lisbon, Lisbon, Portugal e-mail: apsantos@ff.ul.pt

M. Aschner Department of Pediatrics, Vanderbilt University, Nashville, TN, USA

Department of Molecular Pharmacology, Albert Einstein College of Medicine, Bronx, NY, USA e-mail: michael.aschner@vanderbilt.edu; michael.aschner@einstein.yu.edu

#### Abstract

Over the past years there has been considerable progress in our understanding of manganese (Mn)-induced neurotoxicity and its mechanisms. These studies led to changes in Mn safety assessment around the world. However, manganism continues to represent a health concern, especially considering the recent findings linking manganism to Parkinson's disease (PD). Animal models have been invaluable in these investigations. Findings from these studies are discussed in this chapter within the context of Mn-induced neurotoxicity mechanisms and its role in the etiology of parkinsonism.

#### Keywords

Dopamine • Manganese • Neurotoxicity • Parkinson's disease

# 1 Introduction

Over the past several decades, there has been considerable progress in our basic knowledge regarding the mechanisms and factors regulating Mn neurotoxicity. The disorder known as manganism is associated with the preferential accumulation of Mn in the globus pallidus of the basal ganglia, which is generally considered to be the major and initial site of injury. Because the area of the central nervous system (CNS) comprising the basal ganglia is complex and dependent on the precise functioning and balance between several neurotransmitters, it is not surprising that symptoms of manganism overlap with those of Parkinson's disease. The fact that neurological symptoms and onset of Mn toxicity are quite broad and can vary unpredictably probably reflects specific genetic variance of the physiological and biochemical makeup within the basal ganglia; however, there is evidence suggesting that persistent exposure can predispose individuals to acquire dystonic movements associated with Idiopathic Parkinson's disease (IPD). This chapter provides a review on Mn-induced neurotoxicity with specific focus on data using various animal models that provided some key findings in an attempt to unravel mechanisms that lead to these syndromes and what they have in common.

# 2 Manganese Essentiality

Mn is an essential nutrient necessary for a variety of metabolic functions including those involved in normal human development, activation of certain metalloenzymes, energy metabolism, immunological system function, nervous system function, reproductive hormone function, and in antioxidant enzymes that protects cells from damage due to free radicals (ATSDR 2000). In the brain, Mn is an important cofactor for a variety of enzymes, including those involved in neuro-transmitter synthesis and metabolism (Erikson and Aschner 2003). The most important source of Mn is the diet, although there is no consensus regarding safe and adequate levels of this nutrient for various age groups.

Mn deficiencies, although not frequently reported, can result in several biochemical and structural defects in experimental animals (Keen et al. 1999). Thus, taking into account the variety of enzymatic processes which are dependent upon Mn, an inadequate daily supply of the metal may cause a variety of health repercussions, ranging from generalized growth impairment, birth defects, reduced fertility, and impaired bone formation to altered metabolism of lipids, proteins, and carbohydrates (Freeland-Graves and Lin 1991; Keen et al. 1999). Mn is a required cofactor for several enzymes; however, it can be replaced by iron, magnesium, or copper. Notably, the majority of Mn-dependent enzymes known are found in bacteria and plants. The well known Mn-dependent mammalian enzymes include arginase, glutamine synthetase, phosphoenolpyruvate decarboxylase, and manganese superoxide dismutase (MnSOD).

# 3 Manganese Toxicity

# 3.1 Sources of Manganese Exposure

There are many sources of Mn exposure which have been of interest in causing neurotoxicity. Table 1 summarizes the most important sources of Mn intoxication.

# 3.2 Main Routes of Manganese Exposure

# 3.2.1 Pulmonary Tract

Mn is readily absorbed by the pulmonary tract as it lacks the homeostatic control systems that are present in the GI tract. The bioavailability of the inspired Mn depends on several factors, such as the size, mass, and density of the particle, the site of Mn deposition, the composition of Mn within the particulate matter, and how efficiently the metal is solubilized by macrophages within the lung (Roth 2006). It is believed that, at least in part, three characteristics contribute to Mn's absorption via the pulmonary tract: (1) transport into the olfactory or trigeminal presynaptic nerve endings located in the nasal mucosa, (2) transport across the pulmonary epithelial lining fluid, and/or (3) mucociliary elevator for the lung clearance (Roth 2006).

# 3.2.2 Gastrointestinal (GI) Tract

Oral intake of Mn is the major physiological route of absorption, occurring mainly through the consumption of water, fruits, and vegetables (Khan et al. 2008). The mechanism by which Mn is absorbed by this route is not completely understood; however, it is believed that both passive and active transport are involved (Garcia-Aranda et al. 1983). In general, the concentration of Mn in the diet influences the amount absorbed through GI tract as well as the amount which is eliminated in the bile. When dietary Mn levels are high, adaptive changes include

Source	Examples	Nontoxic doses	Reference
Water		<150 µg/L	Ljung and Vahter 2007; Wasserman et al. 2006
Diet	Teas, vegetables, dietary supplements	0.9–10 mg/day	Finley and Davis 1999; ATSDR 2000
Airborne	Welding fumes, MMT <sup>a</sup> , steel foundries, ferromanganese production	<2 µg/day	Lynam et al. 1999; Roth 2009; Zayed et al. 1999
Milk and infant formulas		<105 mg/day	Stastny et al. 1984
Parenteral exposure	TPN <sup>b</sup> , radiological contrast	100 µg/L	Bertinet et al. 2000; Ono et al. 1995
Pesticides	Maneb, Mancozeb		Aschner 2000
Drugs	Methcathinone		Sikk et al. 2011

Table 1 Sources containing Mn that may lead to intoxication

<sup>a</sup>Methylcyclopentadienyl manganese tricarbonyl

<sup>b</sup>Total parenteral nutrition

reduced GI absorption, enhanced Mn liver metabolism, and increased biliary and pancreatic excretion of the metal (Dorman et al. 2001, 2006). Additionally, several factors can influence Mn absorption and retention. For example, dietary intake is influenced by the presence of other trace minerals, phytate, ascorbic acid, and other dietary constituents (Aschner and Gannon 1994; Davidsson et al. 1989).

# 3.3 Manganese Pharmacokinetics

#### 3.3.1 Absorption

The major source of Mn in humans is via dietary ingestion (Au et al. 2009). Approximately 3–5 % of ingested Mn is absorbed through the intestinal wall, and the rest is excreted in feces, representing a precise homeostatic control over Mn absorption. Its uptake is tightly regulated and any excess of ingested Mn is readily excreted via the bile. In contrast, both pulmonary uptake and particulate transport via the olfactory bulb (Verity 1999; Zayed et al. 1999) can lead to Mn deposition in the striatum and cerebellum and inflammation of the nasal epithelium (Roth 2009).

Iron (Fe) has a strong influence on Mn homeostasis as both metals share physiological transporter [transferrin (Tf)], binding and uptake via the Tf transporter and the divalent metal transporter, DMT1/NRAMP2. In rodent models, Fe deficiency is associated with increased Mn absorption across the gastrointestinal tract, as well increased Mn deposition in the brain (Fitsanakis et al. 2008; Freeland-Graves and Lin 1991; Garcia et al. 2007).

## 3.3.2 Transport

The identity of the carrier(s) involved in Mn transport into the brain is still controversial. It is believed that at normal plasma concentrations, Mn enters into the CNS primarily across the capillary endothelium, whereas at high plasma concentrations, transport across the choroid plexus predominates (Murphy et al. 1991). How and in what chemical form Mn is transported across the blood–brain barrier (BBB) has been addressed in series of studies. Mn is absorbed in the GI tract as  $Mn^{2+}$ , is oxidized to  $Mn^{3+}$  by liver and plasma ceruloplasmin, and transported by transferrin (Tf) (Aschner and Gannon 1994; Takeda et al. 1995). Tf-dependent Mn transport across the BBB has been documented (Aschner and Gannon 1994), but the majority of BBB Mn transport occurs via the DMT1.

A critical regulator of brain Mn levels is the divalent metal transporter, DMT-1/NRAMP-2. DMT-1 (also referred to as the DCT or divalent cation transporter) is known to shuttle both Mn and Fe ions in the (+2) valence, as well as other divalent metals. Mutations in *DMT-1* in the rat or mouse result in significantly lower brain levels and uptake of both Mn and Fe (Chua and Morgan 1997; Fleming et al. 1998). Notably, a recent study (Salazar et al. 2008) has shown that DMT-1 contributes to neurodegeneration in an experimental model of PD.

Additional brain Mn transporters include the Mn-citrate transporters (MCT) and the Mn-bicarbonate symporters (Crossgrove et al. 2003). However, the relevance of these genes to Mn transport in vivo is not completely understood. The Mn-bicarbonate symporters, ZIP-8 and ZIP-14, have been identified as members of the solute carrier-39 and are expressed on brain capillaries (He et al. 2006). These symporters utilize a  $HCO_3^-$  gradient as the driving force for Mn uptake across the plasma membrane.

Other possible mechanisms for Mn transport include dopamine transporters (DAT). It is believed that DAT facilitates Mn transport into dopaminergic striatal neurons and that Mn accumulates in the globus pallidus via axonal transport (Anderson et al. 2007). As a result, blockage of the DAT in the striatum attenuates Mn accumulation in striatal neurons and decreases Mn concentrations in the globus pallidus (Anderson et al. 2007). Finally, Mn transport via voltage-regulated channels (Lucaciu et al. 1997), store-operated channels (Riccio et al. 2002), ionotropic glutamate receptor channels (Kannurpatti et al. 2000) (all Ca<sup>2+</sup> channels), and choline transporters (Lockman et al. 2001) has also been described.

## 3.3.3 Distribution

Under basal conditions, brain Mn concentrations are heterogeneous, with the highest values reported in the palladium and putamen in humans and in the hypothalamus in rats (Bird et al. 1984; Dorman et al. 2001). The basal ganglia accumulate the highest Mn levels upon excessive exposure to this metal, although the precise physiological basis for this result remains unclear. However, the high expression of DMT1 in several nuclei of the basal ganglia likely contributes to the propensity of Mn to accumulate in this particular region of the brain (Huang et al. 2004). Similarly, high levels of Mn within the olfactory bulb have been associated with the high expression of DMT1 in this region in rats (Fitsanakis et al. 2008). However, further characterization of the absorption and elimination kinetics, as well as Mn uptake and export pathways are necessary to better understand the basis of differential Mn accumulation

in different brain regions. It is worth noting that recent pharmacokinetic modeling of Mn accumulation in the rat required different diffusion constants between brain regions to account for the asymmetry in Mn levels across the brain (Nong et al. 2008).

It is noteworthy that no information exists on the putative extracellular transport mechanisms of Mn; thus, there is no knowledge regarding conditions that affect the removal of Mn from the brain. However, ferroportin-1 protein expression, the only known iron exporter, is induced by Mn exposure in HEK293T cells (and mouse brain), attenuating Mn accumulation and cytotoxicity and strongly suggesting that ferroportin-1 is a putative Mn exporter (Yin et al. 2010).

## 3.3.4 Excretion

The physiological half-life of Mn in the adult rat and primate brain is approximately 51–74 days (Cotzias et al. 1968; Newland et al. 1987; Takeda et al. 1995). The main excretion mechanism for Mn depends on normal liver function. Indeed, blood Mn concentrations are increased during the active phase of acute hepatitis as well as in post hepatic cirrhosis, and a significant correlation exists between blood Mn and the activities of liver enzymes in patients with hepatitis and cirrhosis (Spahr et al. 1996). Corroborating this hypothesis, magnetic resonance imaging (MRI) has consistently shown signal hyperintensity in the globus pallidus of at least 80 % of cirrhotic patients (Butterworth et al. 1995). Furthermore, direct measurements in pallidal samples obtained from the autopsies of cirrhotic patients revealed several-fold increases of Mn concentrations, and histopathologic evaluations showed Alzheimer's type II astrocytosis (Weissenborn et al. 1995). In fact, hepatic encephalopathy results in cognitive, psychiatric, and motor impairments, which could be related to manganism (Hazell and Butterworth 1999).

# 4 Manganese Neurotoxicity

It has been known for more than 150 years that Mn is a neurotoxic agent. Mn toxicity has been predominantly observed in occupational settings, following the accidental ingestion of large quantities or after chronic inhalation of high levels (Mergler et al. 1994). The brain is particularly susceptible to excess of this metal, but the mechanisms of toxicity are poorly understood. In humans, it has been postulated that there is a spectrum of neurobehavioral and neurophysiological effects associated with Mn toxicity, including both subclinical and clinical symptoms (Mergler et al. 1994).

Mn was first considered to be neurotoxic when five workers employed in grinding the black oxide of Mn developed an unsteady gait and muscular weakness (Couper 1837). Traditionally, Mn neurotoxicity has been most commonly associated with occupations such as Mn mining and smelting, battery manufacturing, and steel production (Santamaria et al. 2007). Nowadays industrial environments, occupations in battery production, welding operations, and mining represent the main sources of Mn exposure and intoxication (Huang et al. 1989; Roth 2009). Furthermore, non-occupational exposures from contaminated drinking water, milk and infant formulas, total parenteral nutrition (TPN) and air as a result of the combustion of the gasoline

additive, methylcyclopentadienyl manganese tricarbonyl (MMT), also present contemporary health concerns (Aschner and Dorman 2006; Aschner et al. 2009; Santamaria et al. 2007).

Mn neurotoxicity, or *locura manganica*, also referred to as manganism, is a neurologic disorder characterized by psychological and neurological abnormalities which resemble Parkinson's disease (PD) (Barbeau 1984; Huang et al. 1989; Mena et al. 1967), Mn also damages brain areas distinct from those that are affected in PD (Calne et al. 1994; Olanow 2004). The similarities between the clinical manifestations of PD and manganism include the presence of generalized bradykinesia and widespread rigidity, while dissimilarities include less frequent resting tremor, more frequent dystonia, symmetry of effects, a particular propensity to fall backward, a characteristic "cock-walk," in which patients walk on their toes with elbows flexed and spine erect (Calne et al. 1994). There are also differences with respect to treatment response – although there may be an initial response to levodopa, the primary treatment option for PD, there is typically a failure to achieve a sustained therapeutic response in patients with manganism (Aschner et al. 2009; Calne et al. 1994). The similarities between the two disorders can be partially explained by the fact that the basal ganglia accumulate most of the excess Mn compared with other brain regions in manganism, and dysfunction in the basal ganglia is involved in PD (Dobson et al. 2004). PD is primarily associated with the loss of dopaminergic neurons within the substantia nigra, allowing the caudate and putamen to become overly active and possibly cause continuous output of excitatory signals to the corticospinal motor control system (Dobson et al. 2004). The substantia nigra is spared in manganism, which is linked to the degeneration of GABAergic neurons within the globus pallidus in pathways postsynaptic to the nigrostriatal system (Pal et al. 1999).

The symptoms observed in Mn-exposed subjects are believed to be caused by the preferential accumulation of Mn in brain areas which are rich in dopaminergic neurons (DAergic) and are largely involved in motor control. Perturbation of striatal DA levels likely explains the biphasic syndrome experienced by manganism patients. Upon Mn intoxication, an initial phase of increased DA production is associated with psychotic episodes commonly observed in psychiatric patients (Aschner et al. 2009; Roth 2009). As Mn poisoning progresses, catecholamine levels decrease, most likely due to the loss of nigrostriatal DAergic neurons and, consequently, the parkinsonian-like symptoms ensue (Benedetto et al. 2009).

## 4.1 Manganese and Mitochondria

Different studies point to the mitochondria as the main cellular targets of Mn toxicity. Nominally, mitochondria have been posited to play a central role in the etiology of both Parkinson's disease (PD) and manganism. Due to transport mechanisms that favor the influx of Mn over its efflux, the metal accumulates in this organelle and thus can readily disturb its function (Gavin et al. 1990). Using

imaging techniques, Morello et al. (2008) reported that prior to any ultrastructural damage to neurons and astrocytes, the largest rate of Mn increase was noted in the mitochondrial cytoplasmic matrix of these cells (Morello et al. 2008).

 $Mn^{2+}$  is sequestered by mitochondria via the Ca<sup>2+</sup> uniporter (Chance 1965). Once inside, Mn<sup>2+</sup> can trigger mitochondrial dysfunction through several mechanisms, primarily involving oxidative stress. Oxidative stress can also occur as a consequence of mitochondrial dysfunction caused by Mn. Studies of postmortem PD brains show damage to the substantia nigra (SN) consistent with generation of reactive oxygen and nitrogen species (ROS, RNS), characterized by lipid peroxidation, protein oxidation, 3-nitrotyrosine formation, DNA oxidation, DNA breaks and a decrease in the activities of the ROS scavenging enzymes glutathione peroxidase and superoxide dismutase (Bowman et al. 2011). Mn can also impair mitochondrial function by inhibiting the mitochondrial electron transfer chain. Zhang et al. (2004) reported that  $Mn^{2+}$  affects the activity of respiratory chain enzymes, inhibiting complexes I-IV and interfering with oxidative phosphorylation (Zhang et al. 2004). Furthermore, it has been shown that Mn can bind to a specific Mg<sup>2+</sup> site on the F1-ATPase, slowing its activity (Smith et al. 1985). Such effects result in a decline of ATP production, leading to increased leakage of electrons and increased  $O^{2-}$  production, thereby causing the eventual collapse of cellular function, which may be followed by cell death (Scholte 1988).

Mn also alters calcium homeostasis in mitochondria by inhibiting its efflux and increasing its levels in the organelle (Gavin et al. 1990). It has been widely reported that elevated calcium levels associated with ROS production result in the opening of the mitochondrial permeability transition pore (MPT), increasing the permeability to protons, ions and solutes (Zoratti and Szabo 1995). As a consequence, there is loss of the inner membrane potential ( $\Delta\psi$ m), leading to mitochondrial swelling, impairment of oxidative phosphorylation and ATP synthesis (Gunter and Pfeiffer 1990). Indeed, Mn has been found to decrease  $\Delta\psi$ m in a concentration dependent manner, indicating that this calcium-dependent process may occur during Mn neurotoxicity (Gunter and Pfeiffer 1990; Milatovic et al. 2007; Yin et al. 2008).

Additionally, apoptotic mechanisms may be also related to Mn neurotoxicity. Calcium-induced MPT opening leads to the activation of the Bcl-2 family of proteins, especially Bax/Bac, culminating with the release of cytochrome c (Cyt c) (Green and Reed 1998; Korsmeyer et al. 2000). Cyt c activates, via the ERK pathway, the cysteine protease caspase-3, the executioner protein, which is responsible for mediating apoptosis, chromatin condensation and DNA fragmentation (Petit et al. 1996). Indeed, Yin et al. (2008) demonstrated that Mn exposure in astrocytes results in ERK and caspase-3 activation. Further, Malecki (2001) observed DNA strand breakage at Mn concentrations as low as 5  $\mu$ M, reinforcing this mechanistic hypothesis.

In addition, the overexpression of wild-type human  $\alpha$ -synuclein in *C. elegans* increases vulnerability to mitochondrial complex I inhibitors, such as rotenone, fenperoximate, pyridaben, and stigmatellin (Weissenborn et al. 1995).  $\alpha$ -Synuclein overexpression has also been recently shown to enhance Mn-induced neurotoxicity via the NF-kB-mediated pathway (Hazell and Butterworth 1999) in a mesencephalic cell line (MES 23.5). Mn also induces the overexpression of  $\alpha$ -synuclein in

PC12 cells via ERK activation (Zlotkin et al. 1995) and chronic exposure to Mn decreases striatal dopamine turnover in human  $\alpha$ -synuclein transgenic mice (Butterworth et al. 1995).

## 4.2 Dopaminergic System

Dopamine (DA) is one of the most prevalent catecholamines within the brain; its signaling helps control locomotion, emotion and neuroendrocrine secretion (Cooper et al. 1996). In this regard, research conducted on Mn toxicity either focuses on the effect on the DA transporter (DAT) or the DA receptor. Postnatal Mn exposure causes a decline in presynaptic DAergic functioning, such as DAT protein expression and DA uptake in the striatum, and a longterm decrease in DA efflux (Huang et al. 2003; McDougall et al. 2008). In adult animal models, there is strong evidence to suggest that exposure to Mn can inhibit DA neurotransmission and deplete striatal DA (Chen et al. 2006; Pal et al. 1999; Reichel et al. 2006), thereby resulting in the observed motor deficits (see review, Guilarte 2010). Importantly, chronic exposure to Mn also decreases striatal DA turnover in human alpha-synuclein transgenic mice (Peneder et al. 2011).

The two main families of DA receptors responsible for DA regulation are D1- and D2-receptors. D1-receptors are G-protein-coupled receptors that stimulate adenylate cyclase activity, while D2-receptors inhibit adenylate cyclase activity (Baldessarini and Tarazi 1996; Hazelwood et al. 2008). A recent study has demonstrated that striatal D2-receptors are not affected by Mn intoxication in monkeys (Eriksson et al. 1992); nevertheless, other studies have shown that the density of D2-receptors is lowered in developing rats treated with Mn (Seth and Chandra 1984).

The direct interaction between Mn and DA is also noteworthy. Both  $Mn^{2+}$  and  $Mn^{3+}$  are known to react with DA via the Fenton's reaction, catalyzing the autooxidation of DA and generating reactive oxygen species (ROS), thereby leading to oxidative damage (Donaldson et al. 1981, 1982; Lloyd 1995; Stokes et al. 1999). Mn-induced DA auto-oxidation relies on various mechanisms involving semiquinone and aminochrome intermediates, L-cysteine or copper (Cu) and NADH facilitation (Graumann et al. 2002; Lloyd 1995). Mn oxidant generation is believed to depend on its oxidation state, and oxidative damage is more pronounced for  $Mn^{3+}$ versus  $Mn^{2+}$  oxidation state (Reaney and Smith 2005).

# 5 Manganese and Parkinson's Disease

Idiopathic Parkinson's disease (IPD) is a progressive neurodegenerative disorder with a slow onset, and compared with the familial forms of the disease, it is associated with advanced age (>55 years old), tremors at rest, bradykinesia, rigidity, and postural instability (Jankovic 2008; Lees et al. 2009; Tolosa et al. 2006).

Molecular imaging is a useful strategy for the diagnosis of PD and has provided extensive evidence that PD patients exhibit decreased levels of presynaptic dopamine neuron terminal markers in the striatum (Felicio et al. 2009). This is consistent with the loss of DA terminals as a result of degeneration of neuronal cell bodies in the substantia nigra pars compacta. Single positron emission tomography studies have shown that PD patients exhibit decreased levels of DA transporters (DATs) and vesicular monoamine transporter type 2(VMAT2) and reduced activity of dopa decarboxylase as measured by the conversion of L-DOPA to dopamine in the striatum (Felicio et al. 2009).

As previously described in this chapter, Mn causes oxidative stress, mitochondrial dysfunction, DAergic degeneration, among other molecular damages, which culminate with an extrapyramidal syndrome called parkinsonism, because of the similarities with PD. While the underlying etiology of the majority of cases of IPD is unknown, there is evidence that exposure to Mn, even at relatively low levels, can result in a spectrum of neurologic impairment including parkinsonism and perhaps IPD itself (Meeker et al. 2007). Manganism like PD is associated with altered neurological dysfunction of the basal ganglia, and because symptoms often overlap between the two, incorrect diagnosis can occur. This is especially true in the latter stages of the disorder when it is feasible that manganism may actually manifest in more Parkinson's-like features in susceptible individuals (Roth 2009).

Characterization of this neurological disorder and investigation of the mechanisms was possible due to research of Mn neurotoxicity in different animal models. A suitable animal model for parkinsonism is histopathologically characterized by the progressive and significant loss of DA neurons and the loss of some non-DA neurons. Moreover, animal models readily mimic the clinical manifestations of the human descriptions.

## 5.1 Studies in Drosophila melanogaster

Drosophila has a relatively simple central nervous system, showing progressive degeneration of DAergic neurons, formation of Lewy bodies, and impairment of climbing activity in PD models (Barone et al. 2011; Feany and Bender 2000). There are about 200 DAergic neurons in the flies; however, they are responsible for modulating a wide range of behaviors ranging from sleep and locomotion to courtship and learning (Van Swinderen and Andretic 2011).

Under chronic Mn exposure, flies show reduced survival, life-span, and reduced locomotor activity, with Mn showing greater toxic effects than iron and copper (Bonilla-Ramirez et al. 2011). Mn accumulates in the head of the flies, and the number of DAergic neuronal clusters labeled with green fluorescent protein (GFP) is reduced upon Mn treatment (Bonilla-Ramirez et al. 2011). This study also showed that Mn-induced DAergic degeneration was associated with impairment in the climbing behavior and mobility shortage analogous to fly PD models.

## 5.2 Studies in Caenorhabditis elegans

The nematode *Caenorhabditis elegans* offers several advantages as a tool for studying neurotoxicity, including its relatively short life-span, lasting 20 days on average, and low costs in growing and maintaining large colonies (Link 2006). Genetic, genomic, and chemical mutant screens are more easily performed in *C. elegans* than in most other experimental species. Compound screening using *C. elegans* is also less time-consuming, due to fast reproduction and a high progeny number (Schmidt et al. 2007). The basic cell biology and biochemistry of nematodes overlap with mammals, including similar ion channels, neurotransmitters (e.g., DA, serotonin, acetylcholine, GABA), vesicular transporters, receptors, and synaptic components, particularly the ones that form the DAergic system (Bargmann 1998). The wiring diagram of the nematode nervous system consists of a defined set of 302 neurons, of which eight are dopaminergic (Gitler et al. 2009). Several worm behaviors are related to DAergic function which also enables to evaluate the effects of the degeneration in this model.

Mn exposure in worms has been associated with morphological alterations, reduced longevity, oxidative stress, and behavioral alterations, such as decreased head trash and body-bend frequencies (Au et al. 2009; Benedetto et al. 2010; Settivari et al. 2009; Wang and Xing 2008). Particularly, DAergic degenerations were observed after acute exposure to this metal (Benedetto et al. 2010; Settivari et al. 2009). Benedetto et al., making use of various genetic strains of C. elegans in conjunction with the fluorescent-tagging of different classes of neuron types, have demonstrated that Mn causes a dose-dependent and specific degeneration of DAergic neurons (Benedetto et al. 2010). This effect was absent in GABAergic, serotonergic, or glutamatergic neurons. A novel finding also showed that this Mn-induced degeneration requires the presence of the reuptake transporter, dopamine transporter 1 (DAT-1), as dat-1 loss-of-function abolished the Mn-induced DAergic loss in the *dat-1*;Pdat-1::GFP worms (Benedetto et al. 2010). In addition, Mn toxicity was reduced by the loss of tyrosine hydroxylase (TH)/CAT-2 function in the double knockout strain, *cat-2(e1112);dat-1(ok157)*, establishing that proper DA synthesis is required for the DAT-1-dependent Mn-induced toxicity. Furthermore, the absence of vesicular monoamine transporter (VMAT2)/CAT-1 in the cat-1(e1111) mutants, where the DA ergic neurons are unable to release DA at the synaptic cleft, resulted in increased tolerance to Mn exposure, indicating that extracellular, but not intracellular DA, is involved in mediating Mn-induced toxicity (Benedetto et al. 2010). Corroborating this finding, loss-of-function of the three DA receptors [dop-1(vs100), dop-2(vs105), dop-3(vs106)] led to increase in extracellular DA, exacerbating Mn toxicity (Benedetto et al. 2010).

The ability of Mn to cause specific DAergic degeneration was linked to the presence in worms of orthologues of the mammalian divalent metal transporter-1 (DMT-1)/NRAMP2 (natural resistance-associated macrophage protein 2). Two of three gene orthologues in worms (*smf-1* and *smf-3*) are mostly localized in the digestive system but are also present in DAergic neurons (Au et al. 2009; Settivari et al. 2009). Decreased susceptibility to Mn and its accumulation in *smf-1* and *smf-3* 

knockout worms demonstrated the involvement of SMF-1 and SMF-3 in Mn uptake in *C. elegans*. Work by Settivari et al. also indicated a role for *smf* in Mn neurotoxicity, demonstrating that genetic knockdown or deletion of *smf-1* and *smf-2* partially inhibits DAergic neuronal death (Settivari et al. 2009).

Pesticides containing Mn are also a source of exposure. Mancozeb (MZ), whose active ingredient is Mn/zinc (Zn)-ethylene-bis-dithiocarbamate (Mn/Zn-EBDC) caused mitochondrial dysfunction and GABAergic and DAergic degeneration in worms (Negga et al. 2011a, b). The use of MZ in the crops has been increasing, yet it has been linked to PD and parkinsonism in humans (Israeli et al. 1983).

# 5.3 Studies in Rodents

Knowledge regarding Mn neurotoxicity and its relation to IPD has been largely derived from rodent studies. A welding fume exposure in rats that corresponded to  $\sim$ 5 years for humans led to alterations in reduced tyrosine hydroxylase levels in the striatum and, more interestingly, to alterations in PD-related proteins expression (Sriram et al. 2010). For instance, there was decrease in the expression of Park5 and Park7. Park5 is involved in the ubiquitin-proteasome system and compromised function or reduced expression of this protein has been associated to IPD (Choi et al. 2004). Park7 mutations account for  $\sim$ 1–2 % of early onset cases of PD, and its reduction may be related to the decrease in mitochondrial integrity (Sriram et al. 2010).

Another PD-related gene that has been investigated is  $\alpha$ -synuclein. The deposition of this small protein in Lewy bodies due to three point mutations in the  $\alpha$ -synuclein gene is a molecular hallmark of familial PD (Ulmer and Bax 2005). It has been reported that Mn can induce overexpression of  $\alpha$ -synuclein in PC12 cells, which reduces cellular viability (Cai et al. 2010). Furthermore, Mn exposure in rat-derived primary cerebellar neuron culture increased levels of  $\alpha$ - synuclein associated to increased oxidative stress (Tong et al. 2009). Furthermore, rat mesencephalic cells expressing  $\alpha$ -synuclein showed higher neurotoxicity to Mn, an effect that occurred through NF- $\kappa$ B pathway activation (Prabhakaran et al. 2011). Combined, these important studies link PD-related genes and manganism, indicating that Mn may increase susceptibility to PD-inherent mutations.

#### 5.4 Studies in Nonhuman Primates

The use of nonhuman primates to study Mn-induced neurotoxicity is of great relevance, considering that they are closely related to humans in relation to genetics, biology, neurotransmission systems, and particularly, behavior. Historically, nonhuman primate studies can be divided into two categories. The studies from the 1960s to the early 1990s used Mn doses that were considerably higher than those used in studies from 1995 to 2008 (for a review, see Guilarte (2010)). Most studies before 1995 used cumulative doses >300 mg/kg body weight. These early studies did not use

neuroimaging techniques because of their more recent availability, with the exception of Newland and Weiss (1992), who administered very low doses of Mn and used T1-weighted magnetic resonance imaging (MRI) (Newland and Weiss 1992).

Administration of exceedingly high doses of Mn (approximately 300 mg/kg/ day) caused loss of postural stability, excitability, hypoactivity, falling, muscular rigidity, tremors, unsteady gait and loss of power in limbs (Eriksson et al. 1992; Neff et al. 1969; Pentschew et al. 1963). Notably, chronic exposure to exceedingly high doses (10 and 15 mg/kg/day) of Mn for >272 days can also cause serious cognitive, behavioral, and motor impairments (Schneider et al. 2006). For instance, videotaped animals showed increased frequency in certain stereotyped behaviors, such as increase in grooming and licking fingers and decrease in climbing in the cage (Schneider et al. 2006).

Decreased DA levels in monkeys brains were usually described (Chandra et al. 1979; Dorman et al. 2006; Newland et al. 1987). However, depending on the exposure design and route of exposure, the areas of decreased dopamine vary from one report to another. Consistently, basal ganglia (especially striatum and globus pallidus) are the brain regions that are more often described as having decreased levels of the neurotransmitter (Bird et al. 1984; Chandra et al. 1979; Eriksson et al. 1987), associated with higher Mn accumulation (Dorman et al. 2006, 2008; Erikson et al. 2007).

# 5.5 Studies in Humans

As pointed out by Perl and Olanow, a very limited number of autopsy studies have been performed on workers who were exposed to Mn or on persons with other related conditions in which Mn would be increased in the brain (Perl and Olanow 2007). Recently, noninvasive imaging techniques to measure Mn effects in the brain have been tested with relative success. For instance, PET (positron emission tomography) scan imaging with 6-[(18)F]fluoro-l-dopa (FDOPA) showed that asymptomatic welders have presynaptic nigrostriatal dopaminergic dysfunction with different anatomical localization from the symptomatic IPD patients analyzed (Criswell et al. 2011). In addition, specific tests showed that these asymptomatic welders were showing motor impairment (Criswell et al. 2011).

Several other authors have proposed that occupational exposure to Mn is a risk factor for parkinsonism (Gorell et al. 1999; Kim et al. 1999; Racette et al. 2001, 2005). This is also verified in reports demonstrating increased frequency of parkinsonism in communities near these occupational sources, becoming an environmental issue. An epidemiological study investigated the risk of Parkinson's disease in the cities of Toronto and Hamilton, Canada, in relationship with industrial emissions of Mn as well as the use of MMT as a gasoline additive (Finkelstein and Jerrett 2007). Subjects were assessed based on residence location to traffic-generated air pollution and neighborhood levels of ambient Mn. Results of this study found an association between ambient levels of airborne Mn in Hamilton and the risk of physician-diagnosed Parkinson's disease. Their statistical modeling

suggests that exposure to ambient Mn correlates with early onset of parkinsonism, consistent with the premise that chronic exposure to Mn adds to the inherent loss of neurons triggered by the normal aging process. A more recent study by Lucchini et al. (2007) evaluated the incidence of Parkinson's disease in pre-Alps Valcamonica valley area adjacent to three ferroalloy plants (Lucchini et al. 2007). These authors indicate that the prevalence of parkinsonian disorders in this area was higher than elsewhere in Italy and Europe having a high crude (296/100,000) and sex- and age-standardized prevalence (407/100,000) when compared to other national and international data (overall prevalence of parkinsonism of 108–257/100,000 in Europe; 157.7/100,000 in Italy) (von Campenhausen et al. 2005).

Mn neurotoxicity is also of great concern. Case reports have been published on patients with hepatic diseases, such as cirrhosis and hepatitis. These patients develop encephalopathy and parkinsonian behavior (Dharmasaroja 2010; Mendieta Zeron et al. 2011; Olanow 2004), associated with reduced liver Mn clearance and increased Mn blood levels, as well as neuronal accumulation, as indicated by the increased t1-weighted signals in MRI (Chalela et al. 2011; Kulisevsky et al. 1993; Rivera-Mancia et al. 2011). Parenteral nutrition has also been linked to encephalopathy and extra-pyramidal symptoms, as the daily doses administered in these preparations are 5–6-fold higher than the required dose (Chalela et al. 2011; Hardy 2009).

#### 6 Conclusion

Over the past few decades considerable progress has been made in understanding Mn neurotoxicity and its mechanisms. Given that Mn accumulates and affects neurons in the basal ganglia, it is not surprising that symptoms of Mn neurotoxicity overlap with those of PD. The studies reviewed herein illustrate that Mn can accelerate the onset of parkinsonism in exposed individuals and that the accumulated Mn in the basal ganglia, which includes both the globus pallidus and the substantia nigra, can lead to DAergic neuronal degeneration and parkinsonism. Furthermore, some subpopulations may display increased vulnerability to Mn-induced neurological deficits becoming most prevalent in late age. The question on whether chronic exposure to Mn directly leads to the development of IPD is difficult to resolve and will require additional studies.

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# Mechanisms of Dopamine Oxidation and Parkinson's Disease

# Juan Segura-Aguilar and Irmgard Paris

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#### Abstract

Dopamine's ability to oxidize to aminochrome explains why this molecule is both an essential neurotransmitter for movement control and a neurotoxic compound that induces toxicity and apoptosis in cell lines. Dopamine spontaneously

J. Segura-Aguilar (🖂)

Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile e-mail: jsegura@med.uchile.cl

I. Paris

Program of Molecular and Clinical Pharmacology, ICBM, Faculty of Medicine, University of Chile, Santiago, Chile

Department of Basic Sciences, Santo Tomas University, Viña del Mar, Chile e-mail: irmgardparis@gmail.com

oxidizes to aminochrome in the presence of oxygen due to the protons on the dopamine hydroxyl groups that are dissociated when dopamine is in the cytosol (pH 7.4). Dopamine oxidation is prevented by vesicular monoaminergic transporter-2 (VMAT-2) that takes up dopamine into the monoaminergic synaptic vesicles where the low pH prevents dopamine oxidation. Dopamine in the cytosol can also be degraded by monoamine oxidase (MAO) and catechol ortho-methyltransferase (COMT) soluble isoform. However, under certain unknown conditions dopamine oxidize to aminochrome, the precursor of neuromelanin, since the neuromelanin pigment is found in the human substantia nigra. Aminochrome participates in two neurotoxic reactions: (i) the oneelectron reduction of aminochrome, which is catalyzed by flavoenzymes that use NADH or NADPH as electron donors. This reaction produces leukoaminochrome o-semiquinone radical, which is extremely reactive with oxygen that autoxidizes, depleting both NADH and O<sub>2</sub> required for ATP synthesis when the flavoenzymes use NADH; and (ii) aminochrome forms adducts with proteins such as alpha-synuclein-inducing and alpha-synuclein-stabilizing neurotoxic protofibrils. In addition, aminochrome inactivate parkin an E3 ubiquitin ligase of proteasomal system, complexes I and III of electron transport chain, tyrosine hydroxylase, actin, and  $\alpha$ - and  $\beta$ -tubulin. Aminochrome is also able to participate in three neuroprotective reactions such as (i) polymerization to neuromelanin, which is a pigment localized in substantia nigra and is present in normal subjects; (ii) aminochrome two-electron reduction to leukoaminochrome catalyzed by DT-diaphorase, which prevents aminochrome neurotoxic reactions; and (iii) glutathione conjugation of aminochrome catalyzed by glutathione S-transferase M2-2. The role of aminochrome in the degeneration of dopaminergic neurons in Parkinson's disease is discussed. Aminochrome may induce the focal neurodegeneration of dopaminergic neurons through mechanisms involving mitochondrial dysfunction, protein aggregation, oxidative stress, and protein degradation dysfunction.

#### Keywords

Alpha synuclein • Aminochrome • Aromatic amino acid decarboxylase • Catechol orto-methyl-transferase • Dopamine • DT-diaphorase (NQO1) • Free radicals • Glutathione-S-transferase M2-2 • Mitochondria complex I • Mono amine oxidase • Oxidation • Parkin • Parkinson's disease • Tyrosine hydroxylase • VMAT-2

List of Abbre	viations
AADC	Aromatic amino acid decarboxylase
COMT	Catechol ortho-methyltransferase
GST M2-2	Glutathione S-transferase M2-2
L-dopa	L-dihydroxyphenylalanine
MAO	Monoamine oxidases
TH	Tyrosine hydroxylase
VMAT-2	Vesicular monoaminergic transporter-2

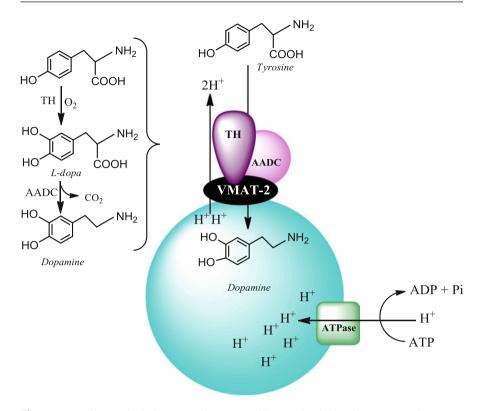
### 1 Dopamine

Dopaminergic neurons play a key role in motor activity. Dopamine is an essential neurotransmitter synthesized from the amino acid tyrosine. It is stored and transported in monoaminergic vesicles, released during neurotransmission, and is taken up by dopamine transporters from the intersynaptic space. Dopamine is synthesized in several steps. The first step involves the formation of L-dihydroxyphenylalanine (L-dopa) from tyrosine by the enzyme tyrosine hydroxylase (TH). The L-dopa is then decarboxylated by aromatic amino acid decarboxylase (AADC) to generate dopamine and CO<sub>2</sub>. Dopamine has two hydroxyl groups that are dissociated in the cytosol at pH 7.4. This dissociation allows the oxidation of dopamine to *o*-quinones. However, dopamine accumulates in monoaminergic vesicles because TH and AADC are associated with the vesicular monoaminergic transporter-2 (VMAT-2). This transporter is present in the membranes of monoaminergic synaptic vesicles and generates a complex with the enzymes (Cartier et al. 2010). Thus, tyrosine is converted to L-dopa by TH and is immediately decarboxylated by ADDC to dopamine. The dopamine is then taken up by VMAT-2 into the monoaminergic synaptic vesicles. The uptake prevents free dopamine from accumulating in the cytosol and the oxidation of dopamine to o-quinone. The dopamine uptake mediated by VAMT-2 is associated with a vesicular ATPase that is translocated into the vessel as ATP is hydrolyzed to ADP, Pi, and one proton  $(H^+)$ . This process creates a proton gradient (Fig. 1). The VMAT-2 transporter exports two protons as one molecule of dopamine is taken into the cell (Chaudhry et al. 2008; Knoth et al. 1981). The pH inside the monoaminergic synaptic vesicles has been estimated to be 2-2.4 pH units lower than the pH in the cytosol (Guillot and Miller 2009). The dopamine inside the monoaminergic synaptic vesicles is stable. It does not oxidize to o-quinone because it is well protonated. The free dopamine in the cytosol participates in three reactions. First, the dopamine can undergo deaminative oxidation to become dihydroxyphenylacetic acid. This reaction is catalyzed by monoamine oxidases. Second, the dopamine catechol structure can be oxidized to become the o-quinone aminochrome. Finally, dopamine can be methylated by ortho methyltransferases (Fig. 1).

# 2 Dopamine Degradation

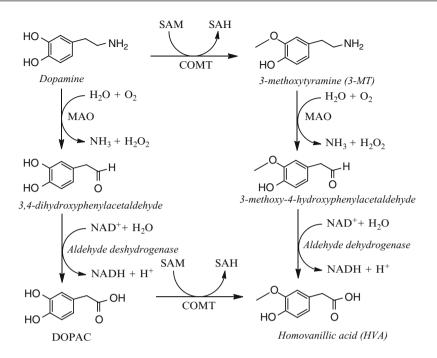
# 2.1 The Oxidative Deamination of Dopamine

The monoamine oxidases (MAO, E.C. 1.4.3.4) degrade excess dopamine in the cytosol by catalyzing the oxidative deamination of the dopamine amino group to 3,4-dihydroxyphenylacetaldehyde. Ammonia and hydrogen peroxide are also produced in this reaction. 3,4-Dihydroxyphenylacetaldehyde is metabolized by aldehyde dehydrogenase to 3,4-dihydroxyphenylacetic acid (DOPAC) as NAD<sup>+</sup> is simultaneously reduced to NADH (Fig. 2). The MAO enzymes



**Fig. 1** Dopamine synthesis from tyrosine catalyzed by tyrosine hydroxylase (*TH*) and aromatic amino acid decarboxylase (*AADC*) and uptake into monoaminergic synaptic vesicles mediated by VMAT-2

are flavoenzymes with multiple isoforms. The isozymes are labeled A and B and share 70 % identity. These enzymes carry sequences specific to FAD (Bach et al. 1988) and are localized to the outer membranes of the mitochondria in neurons, glial cells, and other cell types (Weyler et al. 1990; Shih et al. 1997). MAO-A is mainly found in catecholaminergic neurons, whereas MAO-B is found in serotonergic and histaminergic neurons as well as in astrocytes (Westlund et al. 1988; Saura et al. 1994). MAO-A acts on mainly monoaminergic neurotransmitters such as serotonin, dopamine, noradrenaline, and adrenaline. In contrast, the substrates of MAO-B include trace amines such as tyramine and phenylethylamine (Strolin-Benedetti et al. 1992). The nonselective inhibitors of MAO-A MAO-B include isocarboxazid phenelzine, nialamide, iproniazid, and iproclozide, and tranylcypromine. The selective inhibitors of MAO-A include clorgyline, brofaromine, moclobemide, toloxatone, linezolid, befloxatone, and cimoxatone. The selective inhibitors of MAO-B include selegiline and rasagiline (for review, see Bortolato et al. 2008). The MAO in dopaminergic neurons degrades dopamine to maintain low cytosolic concentrations and



**Fig. 2** Dopamine degradation catalyzed by monoamine oxidase (*MAO*) and catechol *ortho*-methyltransferase (*COMT*)

to prevent dopamine oxidation. However, MAO activity is associated with oxidative stress. This process is a result of hydrogen peroxide formation from the oxidative deamination of dopamine. Hydrogen peroxide is a precursor to the hydroxyl radical. Rasagiline may protect the brain in Parkinson's disease model systems by inhibiting the formation of reactive oxygen species derived from the deaminative oxidation of dopamine catalyzed by MAO-B (Weinreb et al. 2011).

# 2.2 Ortho-Methylation of Dopamine

Dopamine is also degraded by catechol *ortho*-methyltransferase (COMT; EC 2.1.1.6). This enzyme converts dopamine to 3-methoxytyramine, which is then metabolized to 3-methoxy-4-hydroxyphenylacetaldehyde by MAO. Finally, homovanillic acid is formed when aldehyde dehydrogenase catalyzes the oxidation of 3-methoxy-4-hydroxyphenylacetaldehyde (Fig. 2). NADH is also formed in this reaction. Homovanillic acid is also formed by COMT-mediated methylation of the hydroxyl group on DOPAC. Two isoforms of COMT have been discovered: The first is the soluble isoform S-COMT, and the second is the membrane-bound isoform MB-COMT. Both the S- and MB-COMT isoforms are found in microglial

and astroglial cells. COMT is present in pyramidal neurons, cerebellar Purkinje and granular cells, and striatal spiny neurons (Myöhänen et al. 2010). MB-COMT is localized to the cell body, axons, and dendrites of rat cortical neurons. The C-terminal catalytic domain of MB-COMT is in the extracellular space. This domain inactivates the synaptic and extrasynaptic dopamine in presynaptic and postsynaptic neurons (Chen et al. 2011). The downregulation of COMT by estrogen most severely affects the prefrontal cortex and kidneys, where COMT is physiologically important for dopamine metabolism (Schendzielorz et al. 2011). Entacapone has been applied to inhibit COMT in Parkinson's disease treatment. This inhibition prolongs the half-life of L-dopa and allows the delivery of continuous stimulation to the neuronal dopaminergic receptors (Marin and Obeso 2010). A single nucleotide polymorphism (1947) that controls COMT activity has been identified (Tan et al. 2005).

# **3** Dopamine Oxidation to Ortho-Quinones

## 3.1 Dopamine Oxidation to Aminochrome

The protons on the dopamine hydroxyl groups are dissociated when dopamine is in the cytosol (pH 7.4). Dopamine spontaneously oxidizes in the presence of oxygen, even in the absence of metal-ion catalysts (Linert et al. 1996). Dopamine oxidation can produce a dopamine o-semiquinone radical and a superoxide radical (reaction 1). Two dopamine o-semiguinone radicals can react in a disproportionation reaction to produce one molecule of dopamine o-quinone and one molecule of dopamine (reaction 2, Fig. 3). Alternatively, the dopamine o-semiquinone radical can be one-electron oxidized by oxygen to produce dopamine o-quinone and a superoxide radical (reaction 3, Fig. 3). Dopamine oxidation can also be catalyzed by metals. Manganese (III) oxidizes dopamine under aerobic and anaerobic conditions without forming superoxide radicals because manganese (III) catalyzes the formation of both dopamine o-semiquinone and dopamine o-quinone (Segura-Aguilar and Lind 1989). Copper sulfate (II) and iron chloride (III) oxidize dopamine by forming a complex with it (Paris et al. 2001, 2005a, b). Sodium metaperiodate also catalyzes dopamine oxidation (Graham et al. 1978). Further, prostaglandin H synthase, lactoperoxidase, cytochrome P450, xanthine oxidase, and dopamine  $\beta$ -monooxygenase catalyze dopamine oxidation through peroxidase activity (Galzigna et al. 2000; Thompson et al. 2000; Segura-Aguilar 1996; Foppoli et al. 1997; Hastings 1995). Tyrosinase can catalyze the two-electron oxidation of dopamine to dopamine o-quinone. This process does not form dopamine o-semiquinone radicals or consume oxygen (Segura-Aguilar et al. 1998; Jimenez et al. 1984).

Dopamine *o*-quinone is only stable at a pH below 2.0. Thus, this molecule is not stable at physiological pH (Segura-Aguilar and Lind 1989). The amino group of dopamine spontaneously rearranges and undergoes cyclization to become

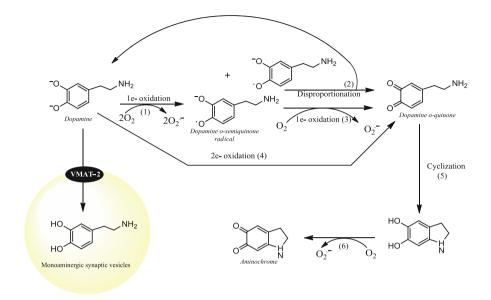


Fig. 3 Dopamine oxidation to aminochrome

leukoaminochrome (reaction 5). Leukoaminochrome undergoes oxidation to become aminochrome (reaction 6). Purified aminochrome (also called dopaminechrome) is stable for approximately 3 h (Paris et al. 2010). This timeframe suggests that aminochrome is the relevant end product of dopamine oxidation in neuromelanin-containing dopaminergic neurons.

Aminochrome formation is dependent on the presence of free cytosolic dopamine that can be oxidized. The efficient uptake of dopamine into monoaminergic vesicles is mediated by VMAT-2, and the degradation of free dopamine by MAO prevents the oxidation of dopamine to aminochrome. However, under certain conditions, free dopamine in the cytosol can oxidize to form aminochrome. The presence of neuromelanin in the substantia nigra of healthy controls suggests that dopamine oxidation occurs in vivo because aminochrome is the precursor of neuromelanin.

# 3.2 Aminochrome Tautomerization

Aminochrome is an unstable molecule that rearranges to 5,6-dihydroxyindole (reaction 1), which is then oxidized to become 5,6-indolequinone (Fig. 4, reaction 2) (Napolitano et al. 2011). However, tyrosinase-catalyzed dopamine oxidation in the absence of metals produces aminochrome that is stable for 3 h after purification (Paris et al. 2010). From a physiological point of view, 3 h is adequate for a molecule to be one- or two-electron reduced by flavoenzymes or to form adducts with proteins. The 5,6-indolequinone is in equilibrium with the quinone methide even though

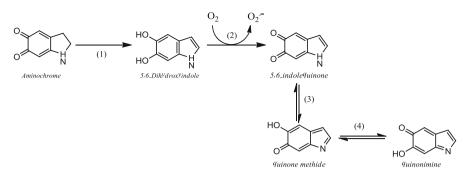


Fig. 4 Aminochrome tautomerization

the presence of transition metal cations favors the formation of the quinone methide (reaction 3). The quinone methide is also in equilibrium with quinonimine (reaction 4, Fig. 4; Pezzella et al. 2007).

## 4 Aminochrome Metabolism

#### 4.1 Aminochrome Polymerization to Neuromelanin

Aminochrome is not completely stable. At high concentrations, it polymerizes to form a dark pigment called neuromelanin (Fig. 5). Neuromelanin formation appears to be a normal process because it is present in dopaminergic neurons of the substantia nigra in healthy subjects without Parkinson's disease. It accumulates with age (Zecca et al. 2002). Neuromelanin appears to have a neuroprotective function through the chelation of metals (Gerlach et al. 2003; Hong and Simon 2007) and the binding of such proteins as alpha-synuclein (Fasano et al. 2006). However, neuromelanin induces apoptosis in cell cultures (Naoi et al. 2008). Interestingly, extracellular neuromelanin also induces microglial activation in the substantia nigra. As a result of this process, superoxide radicals, nitric oxide, hydrogen peroxide, and pro-inflammatory factors are produced, and tyrosine hydroxylase-positive neurons are lost (Zhang et al. 2011). However, neuromelanin is normally stored in double-membrane vacuoles, and the possible neurotoxic effects described by Zhang are avoided.

# 4.2 Aminochrome Adducts with Proteins

Aminochrome forms adducts with proteins that may be involved in the pathogenesis of Parkinson's disease. Alpha-synuclein is hypothesized to cause damage through the generation of neurotoxic protofibrils and has been associated with familial Parkinson's disease. The role of alpha-synuclein in sporadic Parkinson's disease remains an open question. However, aminochrome forms adducts with alpha-synuclein (Fig. 6). This process induces the formation and stabilization of

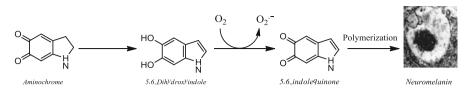


Fig. 5 Aminochrome as precursor of neuromelanin pigment

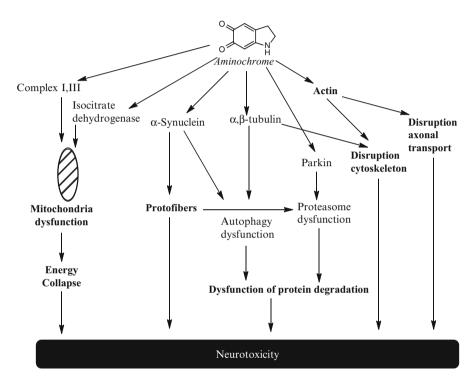


Fig. 6 Formation of aminochrome adducts with proteins

neurotoxic protofibrils that may play a key role in alpha-synuclein neurotoxicity (Conway et al. 2001; Norris et al. 2005). Therefore, aminochrome may be the link between alpha-synuclein and sporadic Parkinson's disease. In familial Parkinson's disease, the formation of neurotoxic protofibrils is dependent on the presence of a specific point mutation that changes alpha-synuclein properties (Polymeropoulos et al. 1997). In sporadic Parkinson's disease, however, the formation of neurotoxic protofibrils is dependent on the formation of neurotoxic protofibrils is dependent on the formation of aminochrome. The aminochrome forms adduct with other proteins such as actin and  $\alpha$ - and  $\beta$ -tubulin and disrupt cytoskeletal architecture (Fig. 6; Paris et al. 2010). Dopamine o-quinone, a transient precursor of aminochrome at physiological pH, forms adducts with and inactivates parkin, a ubiquitin ligase of the proteasomal system (LaVoie et al. 2005).

This result suggests that aminochrome inactivates the proteasomal system, which is consistent with another report (Zafar et al. 2006). Interestingly, a mutation in the parkin gene has also been associated with a familial form of juvenile Parkinson's disease (Kitada et al. 1998). Aminochrome (dopamine o-quinone) forms adduct with the mitochondrial complex I (Van Laar et al. 2009) and induces energy collapse by inactivating mitochondrial electron transport and ATP production. Aminochrome (dopamine *o*-quinone) also forms adducts with and inactivates tyrosine hydroxylase, human dopamine transporter, and tryptophan hydroxylase (Xu et al. 1998; Whitehead et al. 2001; Kuhn and Arthur 1998).

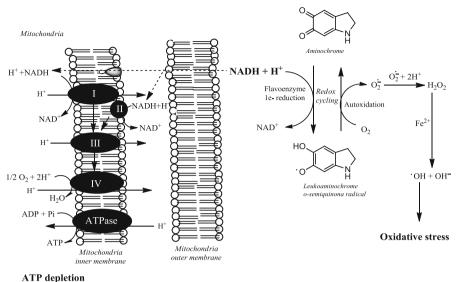
## 4.3 Aminochrome One-Electron Reduction

The aminochrome quinone can be one-electron reduced to become a leukoaminochrome *o*-semiquinone radical. This process is catalyzed by flavoenzymes that transfer electrons from NADH or NADPH to aminochrome. The leukoaminochrome *o*-semiquinone radical is extremely reactive under aerobic conditions. It catalyzes the reduction of oxygen to superoxide and generates a redox cycle between the leukoaminochrome *o*-semiquinone radical and aminochrome (Baez et al. 1995; Segura-Aguilar et al. 1998).

The redox cycling between the leukoaminochrome *o*-semiquinone radical and aminochrome is important for aminochrome neurotoxicity. This process is catalyzed by flavoenzymes and transfers electrons from NADH or NADPH. NADH and oxygen depletion through redox cycling results in energy collapse because these molecules are no longer available for ATP synthesis in the mitochondria (Fig. 7). This effect is potentiated by the formation of superoxide radicals. These radicals spontaneously or enzymatically generate hydrogen peroxide, the precursor of hydroxyl radicals. NADPH can also be used as the electron donor in the reaction catalyzed by flavoenzymes. NADPH is involved in the reaction catalyzed by glutathione reductase that keeps GSH in the reduced state. As a result of NADPH depletion, GSH is unable to exert its antioxidant action, and oxidative stress is thereby increased. The one-electron reduction of aminochrome is neurotoxic in catecholaminergic cells (Paris et al. 2001, 2005a, b, 2009, 2010, 2011; Arriagada et al. 2004; Fuentes et al. 2007; Díaz-Véliz et al. 2008; Muñoz et al. 2012).

#### 4.4 Aminochrome Two-Electron Reduction

DT-diaphorase (EC.1.6.99.2) is a flavoenzyme composed of two subunits and contains two molecules of FAD. This enzyme is unique among flavoenzymes because it catalyzes the two-electron reduction of quinones to hydroquinones. Both NADH and NADPH can be used as electron donators in this reaction. DT-diaphorase is mainly localized in the cytosol, but approximately 5 % of this enzyme is associated with the mitochondria and endoplasmic reticulum. It is found in



All depletion

Fig. 7 One-electron reduction of aminochrome catalyzed by flavoenzymes that use NADH as electron donor

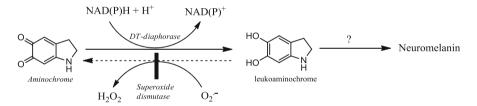
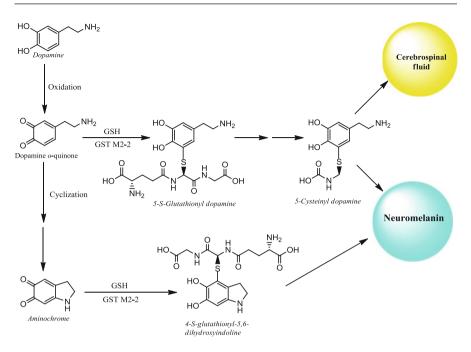


Fig. 8 DT-diaphorase catalyzes the two-electron reduction of aminochrome

different tissues and organs including the brain. DT-diaphorase immunoreactivity co-localizes with tyrosine hydroxylase-like immunoreactivity, suggesting that DT-diaphorase is expressed in the dopaminergic neurons of both the substantia nigra and ventral tegmental area. The dense network of the DT-diaphorase-immunoreactive fibers in the striatum and the dopaminergic innervations were simultaneously lost following a 6-hydroxydopamine lesion. It is important to mention that DT-diaphorase constitutes 97 % of the total cytosolic quinone reductase activity in the rat substantia nigra. DT-diaphorase immunoreactivity has also been found in Bergmann glia, astrocytes, and tanycytes (Schultzberg et al. 1988). DT-diaphorase catalyzes the reduction of aminochrome to leukoaminochrome (Segura-Aguilar and Lind 1989). Leukoaminochrome autoxidizes in the presence of superoxide radicals (Fig. 8). However, the presence of superoxide dismutase



**Fig. 9** Glutathione (*GSH*) conjugation of aminochrome and dopamine *o*-quinone catalyzed by glutathione *S*-transferase M2-2 (*GST M2*-2)

in the cytosol prevents leukoaminochrome autoxidation (Baez et al. 1995). These in vitro results suggest that DT-diaphorase may protect against aminochrome neurotoxicity. Cell culture studies support this idea: DT-diaphorase inhibition with dicoumarol or siRNA induces aminochrome neurotoxicity (Arriagada et al. 2004; Lozano et al. 2010; Paris et al. 2011; Muñoz et al. 2012). DT-diaphorase also prevents the formation of alpha-synuclein protofibrils (Segura-Aguilar et al. 2006; Cardenas et al. 2008) and the cytoskeletal disruption arising from adduct formation with actin and  $\alpha$ - and  $\beta$ -tubulin (Paris et al. 2010).

# 4.5 Aminochrome Conjugation with Glutathione

Aminochrome can be conjugated with glutathione by glutathione *S*-transferase M2-2 (GST M2-2) to 4-*S*-glutathionyl-5,6-dihydroxyindoline (Fig. 9). The 4-*S*-glutathionyl-5,6-dihydroxyindoline molecule is stable in the presence of biological oxidizing agents such as oxygen, superoxide radicals, and hydrogen peroxide (Segura-Aguilar et al. 1997; Baez et al. 1997). This stability suggests that the molecule is a final elimination product. Interestingly, the precursor of aminochrome dopamine *o*-quinone is also conjugated by GST M2-2–5-glutathionyl-dopamine

(Fig. 9). This process prevents the formation of aminochrome (Dagnino-Subiabre et al. 2000). The tripeptide  $\gamma$ -L-Glu-L-Cys-Gly of all glutathione conjugates undergoes glutathione degradation to cysteine conjugation. Therefore, 5-glutathionyl-dopamine is ultimately converted to 5-S-cysteinyl-dopamine (Shen et al. 1996). Interestingly, 5-S-cysteinyl-dopamine has been detected in the cerebrospinal fluid of PD patients and in dopamine-rich brain regions such as the caudate nucleus, putamen, globus pallidus, and substantia nigra as well as in neuromelanin (Cheng et al. 1996, Rosengren et al. 1985, Carstam et al. 1991). GST M2-2 also catalyzes the conjugation of dopa *o*-quinone to 5-glutathionyl dopa. The tripeptide glutathione on 5-glutathionyl dopa undergoes degradation to generate 5-S-cysteinyl-dopa (Dagnino-Subiabre et al. 2000). Melanoma cells produce and release 5-S-cysteinyl-dopa, which is ultimately excreted through urine (Ito et al. 1984). Therefore, the conjugation of glutathione must be protective against aminochrome neurotoxicity.

#### 4.6 Aminochrome and Parkinson's Disease

The discovery that Parkinson's disease patients show a loss of neuromelanincontaining dopaminergic neurons led to intensive research focused on understanding the mechanisms involved in the loss of dopaminergic neurons, which results in motor symptoms. Recently, the presymptomatic and symptomatic phases of Parkinson's disease were subdivided into six stages (Braak et al. 2004). The progressive loss of nigral neurons with Lewy bodies is considered to be an essential neuropathological feature. However, olfactory disturbances, sleep fragmentation, and depression precede the motor symptoms (Wolters and Braak 2006). The disease may initially affect the olfactory bulb, enteric plexus, and motor component of cranial nerve X (stage 1). It may progress to the locus coeruleus, caudal raphe, and magnocellular reticular formation (stage 2). Approximately 5 years before the manifestation of the motor symptoms, the substantia nigra, amygdala central subnucleus, Meynert's nucleus, and pedunculopontine tegmental nucleus are affected (stage 3). Approximately 4 years after the manifestation of motor symptoms, the transentorhinal cortex, CA2 plexus, and intralaminar thalamic nuclei are affected (stage 4). Approximately 10 years after the manifestation of motor symptoms, the prefrontal cortex and tertiary sensory association areas are affected (stage 5). Finally, the secondary and then the primary motor and sensory areas are affected (stage 6) (Hawkes et al. 2010).

Despite the therapeutic efficacy of L-dopa for Parkinson's disease, the role of substantia nigral neurodegeneration (stage 3) in the motor symptoms of the disease remains unclear. The association of the alpha-synuclein (Polymeropoulos et al. 1997) and parkin (Hattori et al. 1998; Abbas et al. 1999) genes with the familial form of Parkinson's disease was important in understanding possible mechanisms underlying the sporadic form of the disease. Several other genes have also been associated with familial Parkinson's disease

(for review, see Corti et al. 2011). Despite the enormous advances in the field, the neurotoxins and mechanisms involved in Parkinson's-related neurodegeneration still remain unclear. However, the mechanisms of dopaminergic neurodegeneration are generally thought to involve mitochondrial dysfunction, alpha-synuclein aggregation in neurotoxic protofibrils, protein degradation dysfunction, oxidative stress, and neuroinflammation (Schapira 2011; Conway et al. 2001; McNaught et al. 2004; Cuervo et al. 2010; Schapira and Jenner 2011).

The proposed timeline for Parkinson's disease (Hawkes et al. 2010) suggests that approximately 5 years elapse between the stage 3 damage to the substantia nigra and the appearance of motor symptoms (stage I in the Hoehn and Yahr scale). The progression of the disease after the manifestation of the motor symptoms is also very slow. It takes approximately 10 years to progress to stage III of the Hoehn and Yahr scale. These data refute the possibility that an exogenous neurotoxin is involved in the idiopathic degeneration of the dopaminergic neurons in the substantia nigra. This possibility is unlikely because MPTP induced severe parkinsonian motor symptoms in 3 days (Williams 1984). It is also important to note that the neuromelanin-containing dopaminergic neurons in the substantia nigra are lost, suggesting that the dopamine in the degenerative cells oxidizes to aminochrome, the precursor of neuromelanin.

We propose that aminochrome is the endogenous neurotoxin involved in the neurodegeneration of the neuromelanin-containing dopaminergic neurons in Parkinson's disease. This compound induces focal neurotoxic events in the years prior to the development of motor symptoms. The following evidence supports this idea: (i) Aminochrome induces mitochondrial dysfunction by forming adducts with complexes I and III in the electron transport chain as well as isocitrate dehydrogenase (Van Laar et al. 2009). The redox cycling induced by aminochrome one-electron reduction depletes NADH, thereby decreasing ATP production in cell culture (Muñoz et al. 2012). (ii) Aminochrome induces the formation and stabilization of alpha-synuclein protofibrils (Norris et al. 2005). (iii) Aminochrome induces protein degradation dysfunction by inactivating parkin (Xu et al. 1998), a ubiquitin ligase involved in the proteasomal system. In addition, alpha-synuclein protofibrils inhibit chaperone-mediated autophagy (Xilouri et al. 2009) and the 26S proteasomal system (Zhang et al. 2008). Aminochrome also induces the aggregation of  $\alpha$ - and  $\beta$ -tubulin through forming adducts with these proteins (Paris et al. 2010). Tubulin aggregation prevents the microtubule formation required for the fusion of autophagocytic vacuoles and lysosomes (Monastyrska et al. 2009). (iv) Finally, aminochrome induces oxidative stress by undergoing one-electron reduction to become the leukoaminochrome o-semiquinone radical (Paris et al. 2001, 2005; Arriagada et al. 2004). Interestingly, aminochrome may play a role in four of the five mechanisms that are generally accepted as being involved in the degeneration of neuromelanin-containing dopaminergic neurons in Parkinson's disease.

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# Neuroinflammation and Parkinson's Disease

## F. Ros-Bernal, J. E. Yuste, E. Tarragón, V. Ortiz, A. Gómez, C. M. Ros, E. Fernández-Villalba, and M. T. Herrero

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F. Ros-Bernal • J.E. Yuste • E. Tarragón • C.M. Ros • M.T. Herrero (🖂)

Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain

e-mail: pacoros@um.es; jeyuste@um.es; ertarr@um.es; carmenmariaros@um.es; mtherrer@um.es; ezquerro@med.uji.es

V. Ortiz • A. Gómez • E. Fernández-Villalba

Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain e-mail: vicortiz@um.es; aurora.gomez@um.es

#### Abstract

Parkinson's disease (PD) is characterized by a slow and progressive degeneration of dopaminergic neurons in the substantia nigra *pars compacta*. To date, despite an intensive research, the cause of the neuronal loss in PD is poorly understood. It has been postulated that neuroinflammatory mechanisms might contribute to the cascade of events leading to this neuronal degeneration. In this review, we describe the evidence for neuroinflammatory processes from postmortem, in vivo and in vitro studies in PD. We further identify the cellular types associated with neuroinflammation that are involved in the degeneration of dopaminergic neurons in animal models and PD patients. Overall, available data support the importance of non-cell-autonomous pathological mechanisms in PD, which are mostly mediated by activated glial and peripheral immune cells. This cellular response to neurodegeneration triggers deleterious events (e.g., oxidative stress and cytokine-receptor-mediated apoptosis), which might eventually lead to dopaminergic cell death and hence disease progression. Novel studies have also demonstrated that genetic markers could be implicated in neuroinflammation and even in the role of glucocorticoids in sustaining activated microglia in inflammatory processes. Finally, based on these evidences, we highlight possible therapeutic strategies aimed at downregulating these inflammatory processes that might be important to slow down the progression of PD.

#### Keywords

Immunomodulatory therapy • Inflammation • Microglia • Neuroinflammation • Parkinson's disease

#### 1 Introduction to Parkinson's Disease

#### 1.1 Overview

Parkinson's disease (PD) is the second most common neurodegenerative disorder in developed countries and the most common neurodegenerative movement disorder (Calabrese 2007; Dorsey et al. 2007; Rochet et al. 2012; Schapira et al. 2009). It is an age-dependent disease with an average age of onset of 55 years of age, approximately 1 % of 65-year-olds are affected, and this rises to 4–5 % in 85-year-olds, while the risk for developing PD increases 5-fold by the age of 70 (Bauso et al. 2012; de Lau et al. 2004; de Rijk et al. 1997; Fahn 2003; Khedr et al. 2012). Two forms of the disease have been characterized: a sporadic form, which generally affects 95 % of all patients and whose etiology is unknown, and a familiar genetic form, which affects 7–10 % of all cases and is linked to mutations in a restricted number of genes (Dauer and Przedborski 2003; Kumar et al. 2012; Martin et al. 2011; Pastor 2012). However, during the last decades, genome-wide association studies (GWAS) have provided evident gains in interpreting the genetic of PD (Nalls et al. 2011; Satake et al. 2009; Simon-Sanchez et al. 2009), and recently, 9 of

PD is characterized by motor symptoms like resting tremor, slowed movement, postural instability, and muscle rigidity (Reichmann 2010); however, there are other symptoms that may include olfactory dysfunctions, urinary problems, constipation, or sleep disruptions and which can halt or even slow down the progression of PD (Erro et al. 2012; Grinberg et al. 2010; Pastor 2012; Peeraully and Tan 2012). The motor symptoms can be treated with dopaminergic drugs; however, the effectiveness diminishes as the severity of the clinical symptoms increases due to progression of the underlying neurodegeneration (Schapira et al. 2009), and moreover, this treatment only improves the symptomatic motor dysfunction.

#### 1.2 Etiology and Molecular Pathways of Neurodegeneration

PD is a neurodegenerative disorder with a complex, multifactorial etiology. In recent years, there has been increasing evidences to support a role for genetic factors in this cause. These evidences have come from twin and familiar studies, where a screening of the most potential susceptibility genes was done. Together with genetic factors, environmental factors may play a role in the disease progress; however, the most accepted theory for PD etiology is that there is a complex interplay between susceptible genes and environmental influences in the causation of PD (Kwok 2010; Soreq et al. 2012).

For non-familiar forms of PD, the dominate view is that multiple causes exist that are decisive in disease initiation and progression; among all of them the age, the genetic predisposition, and the exposition to environmental toxins are especially critical (Nagatsu and Sawada 2006). At the cellular level, three types of cellular alterations may be distinguished in the pathogenesis of PD: the production of reactive oxygen species, dysfunctions in mitochondrial respiratory chain, and an aberrant protein aggregation (Banerjee et al. 2009). Presently, mitochondrial complex I toxins, such as 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP), 1-1'-dimethyl-4-4'-bipyridinium (paraquat), and rotenone, are used for inducing nigrostriatal neuronal degeneration due to dopaminergic (DA) neurons that are specially susceptible to this substances (Meredith et al. 2008).

Following a histological point of view, within the brain the most striking pathological features are the severe loss of dopaminergic neurons in the substantia nigra (SN), the presence of proteinaceous inclusions principally composed of fibrillar  $\alpha$ -synuclein, and ubiquitinated proteins within some remaining nigral neurons called Lewy bodies (LBs) or abnormal neurites called Lewy neurites (LNs) (Lees et al. 2009; Lewy 1912; Trétiakoff 1919). However, nowadays LBs and LNs stained with antibodies to ubiquitin,  $\alpha$ -synuclein, and other biomarkers can be detected in other brain regions apart from the SN, like the locus coeruleus, amygdala, dorsal raphe nucleus, thalamus, and cerebral cortex (Braak et al. 2003; Halliday et al. 2011; McLean et al. 2012). This would explicate why PD debut in patients is associated not with motor dysfunctions but rather with a variety of

non-motor symptoms including those concerning the autonomic nervous system, a subset of which are prodromal and can antecede the motor symptoms by 10 years (Dubow 2007; Grinberg et al. 2010; Kaufmann et al. 2004; Poewe 2007; Tolosa and Pont-Sunyer 2011). According to Braak and Braak's *postmortem* anatomopathological studies, the first areas that are affected by the illness are the dorsal IX/X motor nucleus, medulla oblongata, and pontine tegmentum, and over time, LBs and LNs will affect midbrain regions as SN and finally neocortical regions (Braak et al. 2003). It is not until after reaching specifically the SN *pars compacta* (SNpc) (stage 3 of Braak and Braak classification) that clinical symptoms appear due to the deficits in dopamine release at the level of the striatum (Del Tredici et al. 2002; Picconi et al. 2012).

#### 1.3 Pathogenesis of Parkinson's Disease

Among the environmental toxins, which have been postulated with the development and progression of PD, it has been suggested that pesticides (e.g., paraquat or rotenone), infectious agents (e.g., virus), and heavy metals could have a crucial role (Mizuno et al. 2001; Rohn and Catlin 2011). However, it is not important what insult initially provokes neurodegeneration, since parkinsonism animal models and the function of genes implicated in familial PD form suggest that the main culprits that trigger degeneration in SNpc dopaminergic neurons are the ubiquitinproteasome pathway, mitochondrial dysfunction, and oxidative stress (Elstner et al. 2011; Kanthasamy et al. 2010). While the pathogenic mechanisms that ultimately cause PD are still unclear, because the key molecular and cellular events underlying development of PD are divergent, it is believed that the progressive nature of PD is characterized by the activation of neuroinflammatory pathways that over time trigger neurodegeneration (Dauer and Przedborski 2003; Gao and Hong 2008; Tansey and Goldberg 2010).

#### 2 Neuroinflammation in the Pathogenesis of PD

Glial activation and inflammatory reactions play important roles in DA neuronal death in PD patients and animal models inducing microglial activation, disruption of the blood–brain barrier (BBB), and infiltration of peripheral immune cells (Block and Hong 2005; Choi et al. 2005; Panaro and Cianciulli 2012). However, all these changes are not exclusive of PD, since neuroinflammatory processes are also present in other neurodegenerative diseases such as Alzheimer's disease, Huntington's disease, and amyotrophic lateral sclerosis (Cunningham 2012; Hirsch and Hunot 2009). The purpose of this chapter is to describe the evidences of neuroinflammatory processes associated with the nigrostriatal DA neuronal degeneration not only in the brains of PD patients even more in animal models of parkinsonism. Finally, we will discuss the effectiveness of using specific targets on inflammatory pathways that could relent the progression of PD.

Neuroinflammation is the brain response to infections and injuries associated with most acute or chronic neurodegenerative diseases (Nencini et al. 2003; Schmidt et al. 2005). Neuropathological and neuroradiological studies have shown that prior to significant loss of neurons, a neuroinflammatory process characterized by an increase in glial activity often occurs (Frank-Cannon et al. 2009; Fuhrmann et al. 2010; Ratai et al. 2011). The dogma that the nervous and immune systems were isolated with litter interaction except during disease and/or trauma was based on the evidences that immune responses were blunted in the central nervous system (CNS) and that the blood-brain barrier (BBB) prevented infiltration of immune cells and molecules into the CNS (Prendergast and Anderton 2009). However, recently, the convergence of unexpected results from diverse fields has demonstrated a bidirectional communication between these systems. Firstly, it has been postulated for many years that the BBB was an impermeable barrier to immune cells and most diffusible factors produced in the periphery. However, last studies have clarified that, in response to insult, the permeability of BBB may increase or become dysregulated and also that cytokines produced by glial cells in the brain can cross the compromised BBB and activate immune cells (Deverman and Patterson 2009; Prat et al. 2001). In response to cytokine activity, blood-derived monocytes will migrate through the compromised BBB into the CNS and aid microglia in causing neural inflammation, degeneration, and cell death (Hakimi et al. 2011). Moreover, immune cell infiltration into the CNS is more common than previously believed. Indeed, many neurological diseases, including Alzheimer's and Parkinson's disease, are accompanied by neuroinflammation, and antiinflammatory drugs reduce the risk for Alzheimer's and Parkinson's disease (Lucin and Wyss-Coray 2009). Secondly, there is now growing evidence that neural-immune cross talk may even occur in non-disease conditions, including in the healthy brain, where the permeability of the BBB can be regulated under normal conditions (Chung et al. 2010).

The terms neuroinflammation and neurodegeneration are intimately linked since neuroinflammation consists of an innate immune response which involves the activation of microglial cells which produces several products like cytokines, chemokines, complement cascade proteins, prostaglandins, and reactive oxygen species in response to a peripheral or central insult (Chung et al. 2010), and the term "neurodegenerative" disease, as PD and AD are, is based on the feature that it is primarily the death and dysfunction of neurons that cause disability, being, the activation of various components of the innate immune system, recognized as a feature of these "neurodegenerative" diseases (Tansey and Goldberg 2010). This association is very important under the clinical point of view because if the immune system, as it is supposed, plays any active role in a neurodegenerative disease, for better or for worse, then modifying it – enhancing favorable behaviors or suppressing harmful ones – may be capable of slowing the progression of the diseases is still controversial, since neuroinflammatory response may exert protective but also

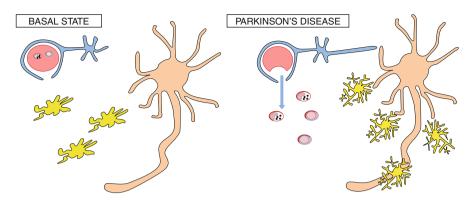
detrimental effects, the reason why it is suggested to be a "double-edged sword" (McGeer and McGeer 2004; Wyss-Coray and Mucke 2002).

#### 2.2 Cell Types Involved in Neuroinflammation

In innate immunity, peripheral effectors are myeloid-derived blood monocytes that additionally give rise to macrophages and a specific type of antigen-presenting cells (APC) that are dendritic cells (DC) (Kurant 2011). The role of these macrophages/ APC is performed by the brain microglia in the CNS, being the primary effectors of immune surveillance and self-renewing and arising from fetal myeloid progenitors (Eglitis and Mezey 1997). Microglia play a homeostatic role in the CNS and respond to environmental stresses and immunologic challenges by removing dying cells (Nakamura 2002), entering a state of chronic activation in which they release toxic factors resulting on ongoing damage to surrounding tissue (Ransohoff and Perry 2009). Although the key molecular and cellular events underlying development of neurodegenerative diseases are clearly divergent among them, one common way in which these divergent molecular or cellular events (e.g., aberrant protein misfoldings, peroxidations, mutations, aggregation) may all contribute over time to neuronal death is activating resident microglial populations in specific brain areas. In the case of genetic mutations or prolonged/repeated environmental exposures, where the initial stimulus that triggered microglial activation is not silenced/eliminated, a self-sustaining cycle of neuroinflammation can ensue, and such a chronic inflammatory environment is presumptive to cause neuronal dysfunctions and death of compromised neuronal populations.

Astrocytes, another member of the glial cell family, contribute to virtually every aspect of brain function, including ionic homeostasis, energy metabolism, and synaptic signalling, and play an important role in the stabilization of the permeability of the BBB (Zlokovic 2008). The varied and important roles of astrocytes have evolved to allow increasingly complex nervous systems to operate efficiently and with high fidelity. For example, astrocytes figure prominently in glutamatergic synaptic transmission, a basic event of brain function: high-affinity glutamate uptake into astrocytes meliorates the temporal and spatial fidelity of glutamatergic signalling, and astrocytes afterward move glutamine back to neurons for the synthesis of more glutamate (Hertz and Zielke 2004; Uwechue et al. 2012). Recent studies have demonstrated that the important and dynamic contributions of astrocytes to normal brain function demand that neuron and astrocyte relationship must be seen as an equal partner relation where this harmonious collaboration will produce a desired function (Ransom and Ransom 2012). So, it has to be discarded as the historical view of astrocytes as simple "scaffolding" cells; nowadays the most accurate view is to denominate astrocytes as "partner cells" (Ransom and Ransom 2012).

During past decades the role of peripheral immune cells in neuroinflammation was very controversial. Besides this controversy, increased accumulation of activated microglia and astrocytes and leukocytes adhering to postcapillary venules are



**Fig. 1** Neuroinflammation triggers the activation of the peripheral immune system. Under neuroinflammatory conditions, there is a modification of the local microenvironment through early microglial activation that allows peripheral immune cell infiltration. Contributing to neuroinflammation, the presence of newly created vessels that have not likely developed the restrictive properties of the blood–brain barrier, failing to protect the parenchyma from peripheral immune cells and inflammatory factors in the peripheral circulation, has been shown

observed in the affected brain areas in neurodegenerative diseases, suggesting the presence of an ongoing inflammatory process (Casoli et al. 2010; Chung et al. 2010). As neuroinflammation triggers the activation of peripheral immune system, several studies implicated subsets of T cells taking part of the neurodegenerative processes. The mechanism related with this specific recruitment of T cells is still unknown, but it might involve a modification of the local microenvironment through early microglial activation and innate neuroinflammatory processes (Chung et al. 2010) (Fig. 1).

#### 2.3 Evidence for a Role of Neuroinflammation in PD

#### 2.3.1 Role of Microglia in PD

Data from postmortem studies provided the first evidence for a role of neuroinflammation in PD, since a massive microglial activity was detected in the brain of PD patients (McGeer et al. 1988b). Moreover, in postmortem investigations of humans exposed to MPTP, activated microglia have been detected 16 years after the last drug exposure (Langston et al. 1999). Since McGeer's initial observation (McGeer et al. 1988b), activated microglia have been identified in the putamen, hippocampus, transentorhinal cortex, cingulated cortex, and temporal cortex of PD brains (Imamura et al. 2003). However, since these are end-stage cases, whether this activation was a general consequence of neuronal death or the reflection of the active participation of microglia in the neurodegenerative process remains unclear (Graeber and Streit 2010).

In order to elucidate the role of microglia activation in PD, several groups have studied several animal models using MPTP, rotenone, and 6-OHDA that exhibit levels of microglial activation similar to what is found in PD (Arai et al. 1999;

Barcia et al. 2003; Carvey et al. 2005; Cicchetti et al. 2002; Gao et al. 2002). However, recent studies have demonstrated that in the case of paraquat and rotenone, microglial activation are likely non-cell autonomous and may occur as a result of dopaminergic neuron damage or by factors released by neurons and/or other cells (Klintworth et al. 2009). Similar to what has been described in MPTP-intoxicated humans, an activated microglia and dopaminergic cell loss was also found in the SN of primates years after they were treated with MPTP (Arai et al. 1999; McGeer et al. 2003), even in those parkinsonian animals never treated neither with levodopa nor with dopaminergic agonists (Barcia et al. 2004), suggesting that microglia plays an active role in the pathology of PD and may indeed perpetuate the degeneration of dopaminergic neurons once activated.

Moreover, the presence of microglial activation participating in neuroinflammatory processes has also been confirmed on a molecular basis. Levels of proinflammatory mediators, including TNF $\alpha$ , IL-1 $\beta$ , IL-6, reactive oxygen species (ROS), and eicosanoids, are elevated in the brains and peripheral blood mononuclear cells (PBMC) of PD patients (McGeer et al. 1988b; Mogi et al. 1994; Nagatsu et al. 2000; Nagatsu and Sawada 2005).

Oxidative stress is caused by an imbalance between the production and destruction of reactive oxygen species (ROS) and reactive nitrogen species (RNS). The production of ROS/RNS under neuropathological conditions would be modulated by microglial NADPH oxidase and nitric oxidase synthase (NOS) (Paramo et al. 2010). Nitrite (as an indicator for nitric oxide free radicals) in the cerebrospinal fluid, as well as increased expression of inducible nitric oxide synthase (iNOS) within the SN, was also found in PD patients (Qureshi et al. 1995). Moreover, an association between Parkinson's disease and polymorphisms in the nNOS and iNOS genes has been described (Levecque et al. 2003). In the MPTP model of parkinsonism, NADPH oxidase and inducible nitric oxidase synthase (iNOS) are known to be major sources of ROS/RNS production where they mediate DA neuronal death in SN (Liberatore et al. 1999; Wu et al. 2002, 2003). Moreover, other in vivo and in vitro studies have demonstrated that microglia-derived NADPH oxidase/iNOS play a crucial role in the production of ROS/RNS and conduce to DA neuronal death following the oxidative stress pathway (Qin et al. 2005; Rodriguez-Pallares et al. 2007). Consequently, the vast majority of therapeutic attention directed at inflammation within the CNS in patients with PD has been directed toward microglial cell activity.

Numerous investigations have proposed a deleterious role of microglial activation in PD based on the vulnerability of dopaminergic neurons to various microgliaderived proinflammatory cytokines (De Lella Ezcurra et al. 2010; Ferrari et al. 2006; Stone et al. 2009), while  $\alpha$ -synuclein can directly induce activation of microglia and release of proinflammatory cytokines, leading to dopaminergic neurodegeneration (Hirsch and Hunot 2009; Zhang et al. 2005). Using animal models, it has been shown that  $\alpha$ -synuclein overexpression triggers an early microglial activation in the SN as well as elicits an increased expression of proinflammatory molecules such as IL-1 $\beta$  and TNF $\alpha$ , prior to dopaminergic neurodegeneration (Kachroo and Schwarzschild 2012; Mosharov et al. 2009; Thiruchelvam et al. 2004). In addition,  $\alpha$ -synuclein is released from cells in vitro and is capable of directly activating microglia (Jin et al. 2007; Maguire-Zeiss 2008; Zhang et al. 2005). Taken together, these observations place  $\alpha$ -synuclein and microglial activation together as early players contributing to neuroinflammation that may mediate ulterior dopaminergic neurodegeneration (Gao et al. 2011).

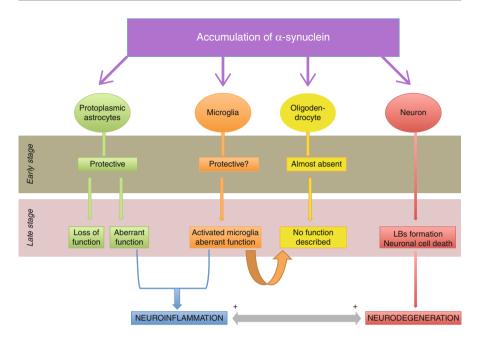
However, it seems that the plasticity of microglia must be considered with regard to their contribution in PD, and their role (whether beneficial or detrimental) may depend on the stimuli present and the stage of disease (Li et al. 2007; Michelucci et al. 2009; Sanchez-Guajardo et al. 2010).

One approach that can connect microglial activation with PD pathogenesis involves in vivo imaging. Positron emission tomography (PET) imaging in combination with the isoquinoline <sup>11</sup>C-PK11195 BP<sub>ND</sub>, which is a ligand of the peripheral binding site of benzodiazepine, is used to track microglial activation (Gerhard et al. 2006a; Ouchi et al. 2005). Although there was a controversy about the efficiency of this marker as a microglial detector, based on discrepancies between in vitro and in vivo experiments, nowadays it is accepted that PK11195 binds to the peripheral binding site of benzodiazepine on activated microglia and macrophages. A significant increase in this PK1195 marker, reflecting microglial activation, was reported in the midbrain and putamen of PD patients when compared to controls, and this activation positively correlated with the motor severity of PD (Bartels et al. 2010; Ouchi et al. 2005). These findings demonstrated the crucial pathogenic importance of early microglial activation for the progression of the disease which suggested that an early introduction of neuroprotective drug to slow down or inhibit microglial activation could be a suitable therapeutic target in PD. Additionally, PD patients exhibited significantly increased <sup>11</sup>C-PK11195 BP<sub>ND</sub> in the basal ganglia, pons, and frontal and temporal cortical regions (Gerhard et al. 2006a). Gerhard and colleagues demonstrated that the increased microglial activation remained unchanged for 2 years, while the patients deteriorated clinically during this period (Gerhard et al. 2006a), similar to what had been described in MPTP-induced parkinsonian monkeys (Arai et al. 1999; Barcia et al. 2004). In relation with this sustained microglial activation, it has been recently described that IFN- $\gamma$  signalling, associated with TNF- $\alpha$  expression, has a crucial and cell-specific role in stimulating and maintaining glial cell activation in parkinsonism (Barcia et al. 2011).

In sum, PET imaging and in vivo and in vitro studies support the hypothesis that microglial activation occurs early in PD where they remain activated for longer periods and possibly drive progression of the disease (Gerhard et al. 2006a, b) (Fig. 2).

#### 2.3.2 Role of Astroglia and Oligodendroglia in PD

Astrocytes, the major glial components of the CNS, constitute up to 20–50 % of brain volume and provide the optimal microenvironment for neuronal function by exerting active control on the cerebral blood flow (Gordon et al. 2008; Mulligan and MacVicar 2004). Although the presence of reactive astrocytes is evident in the most affected brain areas of PD patients having a role in the progression of the disease (Forno et al. 1992; Teismann and Schulz 2004), the specific role of this cell type is still unclear. It was previously described that one of the main functions of astrocytes is to secrete neurotrophic factors in SN for inducing the survival of dopaminergic



**Fig. 2** Effect of  $\alpha$ -synuclein accumulation on neuroinflammation associated with PD. Associated with PD, there is an accumulation of  $\alpha$ -synuclein at different levels. During the early stages of neurodegeneration, astrocytes may quickly respond to  $\alpha$ -synuclein released from degenerating neurons to protect against neurodegeneration. During the long time course of neurodegeneration, the activated astrocytes may lose their protective properties or might be aberrantly activated, leading to stimulation of neuroinflammation. On the other hand, microglia may be as well protective during the early stages of neurodegeneration, but at late stages of the disease, it would be aberrantly activated, contributing to neuroinflammation and perpetuate neurodegeneration. Concerning oligo-dendrocytes, the presence of  $\alpha$ -synuclein deposits in oligodendroglial cells has been described, but its function is still not clear

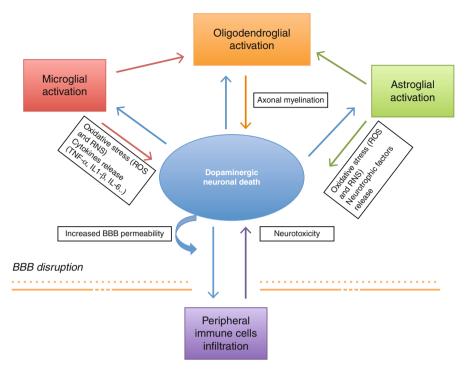
neurons (McGeer and McGeer 2008). Although the relationship between reactive astrocytes and the release of neurotrophic factors is ambiguous, based on previous studies it was concluded that the decreased levels of astrocyte-derived neurotrophic factors are directly responsible for dopaminergic neurodegeneration in PD (Knott et al. 2002; Saavedra et al. 2006; Sandhu et al. 2009).

As it was previously described in this chapter, Lewy pathology follows a progressive pattern beginning in the anterior olfactory structures, dorsal motor nucleus of the vagal nerve, and portions of the enteric nervous system and evolving in caudo-rostral manner within the CNS (Braak and Del Tredici 2008). Unexpectedly, as PD progressed,  $\alpha$ -synuclein-positive inclusions were also detected in astrocytes that were not known to produce  $\alpha$ -synuclein (Wakabayashi et al. 2000). However, recent works have demonstrated, in cultured cells and  $\alpha$ -synuclein overexpressed animal models, that  $\alpha$ -synuclein proteins released from neuronal cells are taken up by astrocytes through endocytosis and form inclusion bodies, and this abnormal inclusions incite changes in astrocytic gene expression profiles reflective of a proinflammatory response (Lee et al. 2010; Tousi et al. 2012). Finally, it was recently described that of the two main types of astrocytes, fibrous and protoplasmic, only protoplasmic astrocytes are involved in PD, where they become nonreactive and accumulate  $\alpha$ -synuclein (Halliday and Stevens 2011). In sum, glial cells are responsible for the progression of PD and play an important role in initiating the early tissue response. In particular, early dysfunction and  $\alpha$ -synuclein accumulation in astrocytes causes recruitment of phagocytic microglia that attack selected neurons in restricted brain regions causing the clinical symptoms of PD (Halliday and Stevens 2011) (Fig. 2).

In relation with oligodendrocytes, although neuroinflammation associated with PD has been attributed to activated microglia and astrocytes in both SN and striatum, the role of oligodendrocytes is unknown. Despite the presence of  $\alpha$ -synucleinpositive filamentous inclusions in oligodendrocytes in the midbrain of PD patients, which was almost absent in PD patients with mild nigral degeneration (Khedr et al. 2012; Wakabayashi et al. 2000), the correlation between oligodendrocytes and parkinsonism has been barely explored (McGeer and McGeer 2008; Takagi et al. 2007). Recently our group has described in the MPTP model of parkinsonism that the loss of dopaminergic neurons and of their axons projecting to the striatum, in MPTPinduced parkinsonism in both mice and macaques, is accompanied by a prominent oligodendrogliosis along the nigrostriatal pathway, which significantly presents a negative correlation with striatal dopaminergic innervation (Annese et al. 2012). This exacerbation in oligodendroglial response could be induced by the cytokines released by both microglia and astrocytes involved in neuroinflammation (Arai et al. 1999: McGeer and McGeer 2008) (Fig. 3). However, the function and mechanisms triggered by oligodendrocytes and the importance of these  $\alpha$ -synuclein deposits must be further investigated in both experimental models of parkinsonism and PD.

#### 2.3.3 Role of T Cells in Mediating Neuroinflammation-Induced PD

A role of T cells in neuroinflammation associated with neurodegenerative diseases had been several times denied based on the idea that the CNS has traditionally been considered "immune privileged" due to the presence of innate microglia and the protection of the BBB, but recently numerous evidences have discarded this hypothesis. In relation with PD, a role for an adaptive immune response in the etiology of PD has been also demonstrated. For example, the presence of T lymphocytes in the midbrain of PD patients suggests that the potential role of infiltrated peripheral cells is related to PD pathogenesis (McGeer et al. 1988a). Even more, T cell infiltration has been found in CNS tissues of PD patients (Brochard et al. 2009; Miklossy et al. 2006), and a markedly increase in the density of IFN-y-positive cells has been detected in brains of PD patients (Hunot et al. 1999). Moreover, there are reports indicating that nitrated  $\alpha$ -synuclein activates peripheral leukocytes in draining lymphoid tissue and mediated adaptive immune responses in potentiating microglial activation and exacerbating neuronal death (Benner et al. 2008). However, whether this inflammation is the consequence or the cause of neuronal injury is still unclear. Recently, it was reported that CD4<sup>+</sup> T cells are cytotoxic in a MPTP parkinsonism model; therefore, invading CD4<sup>+</sup>



**Fig. 3** Processes implicated in neuroinflammatory pathway associated with PD. Dopaminergic cell death induces microglial activation, a reactive astrocytic response, and also blood-brain barrier (BBB) disruption. Through the "hole" in the BBB, peripheral immune cells are allowed to infiltrate. This exacerbated neuroinflammation inducing increased neuronal death through the release of ROS/RNS oxidative products, cytokines, and neurotrophic factors

T cell-mediated immune response contributes to DA neurodegeneration through the Fas/FasL pathway (Brochard et al. 2009). These studies innovatively implicate the adaptive immune system, similar to the innate immune system, not only in the response to but also in an active participation in the pathogenesis of PD. However, more work needs to be done to determine if and how they will serve as a potential target for therapy in PD (Fig. 3).

#### 2.4 Genetic Markers in PD

Numerous monogenic genes have been associated with familial forms of PD (Satake et al. 2009; Simon-Sanchez et al. 2009; Tan and Skipper 2007; Tan et al. 2007), but this accounts for only 10 % of the PD population, a reason why the relative contribution of genes and environmental factors in these cases of sporadic PD is still unclear (Tan and Skipper 2007). In the last years, a few common genetic risk variants could be associated with PD, but this is far to explain the etiology of all the cases (Hill-Burns et al. 2011; Tan et al. 2007). However, recent genome-wide

association studies have identified genetic markers linked to neuroinflammation, providing an open window to understand the pathophysiologic mechanism of PD (Hamza et al. 2010; Nalls et al. 2011; Satake et al. 2009; Simon-Sanchez et al. 2009). In the last decade, various studies suggested a possible influence of HLA alleles (HLA-DR and HLA-DQ) in PD (Hill-Burns et al. 2011; Montgomery et al. 2010; Stranger et al. 2007).

This hypothesis was demonstrated in anatomopathological studies, where PD brains exhibited upregulation of DR antigens and presence of DR-positive reactive microglia (McGeer and McGeer 2008), and in a genome-wide association study, which demonstrated a specially significant correlation between HLA-DR allele and late-onset PD (Hamza et al. 2010; Hill-Burns et al. 2011; Kumar et al. 2012). On the other hand, other studies have analyzed the relationship between a given genetic polymorphism in neuroinflammation-associated genes and the risk of PD. The most consistent genetic evidence implicates TNF in the initiation and progression of PD; a single polymorphism in the TNF promoter resulted in a higher TNF production especially overrepresented in early-onset sporadic PD (Nishimura et al. 2001; Wahner et al. 2007; Wu et al. 2007). On the contrary, TNF polymorphisms affecting its receptor 1 have been found to be significantly decreased in PD patients (Kruger et al. 2000). Other studies on single-nucleotide polymorphisms in the IL-1 $\alpha$  and IL-1 $\beta$  families also corroborate the genetic role in the PD-associated neuroinflammation (Hirsch and Hunot 2009).

This genetic association between inflammatory genetic markers and PD suggests a strong influence of genes on the neuroinflammation process linked to PD and offers new targets for drug development and pharmacogenetics.

#### 2.5 Glucocorticoids and PD

The human being is born with the ability to respond to stress, and the physiological stress response begins when the brain detects a homeostatic challenge and activates the sympathetic nervous system, which releases catecholamines, followed by the slower activation of the hypothalamic-pituitary-adrenal (HPA) axis: hypothalamic secretion of corticotropin-releasing hormone into the pituitary portal circulation activates pituitary secretion of adrenocorticotropic hormone, which then induces the glucocorticoids (GCs) that are secreted by the adrenals (Papadimitriou and Priftis 2009; Sorrells et al. 2009). Interestingly, the functioning of the HPA axis exhibits an enormous individual variability that aids to justify the considerable individual differences in vulnerability to stress-related disease, including psychiatric disorders and neurodegenerative diseases.

GC-GR responses to neuronal injury are complex, for instance, an excess of GCs (such as occurs in chronic stress) was found to aggravate neuronal injury in experimental ischemia (Sorrells et al. 2009), while an important neuronal survival effect of GR was presented in an acute lipopolysaccharide (LPS)-induced inflammatory lesion model (Nadeau and Rivest 2003). Several reports have proposed that the GC-GR responses might be crucially linked to PD pathogenesis. Thus, chronically high levels of GCs were shown to exacerbate motor deficits in 6-hydroxy-DA-treated rats (Smith et al. 2008). This experimental observation corroborates clinical data showing that stress can trigger or worsen motor symptoms in PD patients (Smith et al. 2002). The involvement of GC-GR in DN survival after MPTP intoxication was suggested previously in adrenalectomized mice (Sugama et al. 2009) as well as in transgenic mice harboring antisense GR (Morale et al. 2004).

Recent studies by our group, linking neuroinflammation in PD pathogenesis and the mechanistic actions of GR, exposed that GR is associated with the p65 subunit of NF- $\kappa$ B in microglia nuclei and modulates the transactivation potential of NF- $\kappa$ B (Ros-Bernal et al. 2011). GC-GR system is altered in PD and this may adversely affect DN survival (Barcia et al. 2009). GR dysfunction in PD may result in a chronic inflammatory reaction, and further in-depth work on its glial actions might open innovative therapeutic perspectives.

#### 3 Anti-inflammatory Therapy in PD

The hypothesis that neuroinflammation is extremely implicated in PD progression has become more and more convincing due to the accumulation of preclinical studies, imaging studies, and *postmortem* analysis studies that strongly implicate inflammatory processes in the progressive degeneration of the nigrostriatal pathway. Up to now, it has to be determined if anti-inflammatory therapy in humans could induce any beneficial alteration in PD progression.

Previous clinical trials with anti-inflammatory compounds have failed probably because of many reasons such as the advance stage of the patients enrolled, the dose regimen, or the wrong chosen compound. According to this, minocycline, which was successful in a phase II clinical trial, attenuating nigrostriatal DA neurodegeneration in an animal model (Du et al. 2001; Peng et al. 2006; Ravina et al. 2006), has been proved in a phase II clinical trial with controversial effects (Dodel et al. 2010). Therefore, we need more clinical studies before dismissing the effectiveness of long-term anti-inflammatory compounds. However, taking into account that the innate inflammatory response is not always detrimental, even beneficial (Wyss-Coray 2006), maybe the long-term inhibition of inflammatory response is detrimental for PD treatment (Kanthasamy et al. 2010; Wyss-Coray and Mucke 2002). Instead, more selective targets of inflammatory mediators and participants in cell-death pathways could be considered in future therapies to prevent neuronal loss in PD.

Recent studies have demonstrated that specific drug treatment does appear to reduce susceptibility to PD and immunomodulatory therapies are currently a major area in PD research (Appel et al. 2010; Li et al. 2011). Based on peripheral immune cell activation, immunomodulatory therapeutic strategies may develop vaccines for antigens to trigger cell-mediated anti-inflammatory responses (Benner et al. 2008; Brochard et al. 2009) as well as, once it is clear that BBB is altered in PD, to hamper the migration of immune cells across this BBB (Tan and Skipper 2007). Moreover, these immunomodulatory strategies may also be of

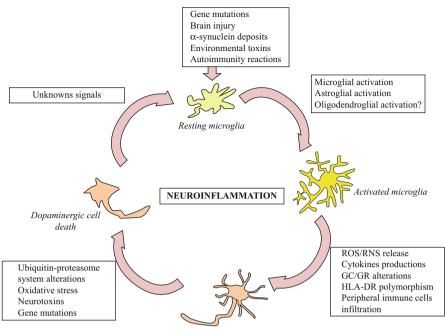
interest for cell transplantation therapies in order to induce a longer survival of grafted DNs which are affected by such an immune-related hostile microenvironment (Satake et al. 2009).

#### 4 Conclusion

Neuroinflammation is widely considered to participate in the progression of PD (Table 1). Microglia-derived inflammatory factors such as TNF- $\alpha$  and IL-1 $\beta$  trigger harmful downstream signalling pathways and accelerate neuronal death in the SN. Activated microglia also increase the production of ROS that impose oxidative stress on the intra-neuronal antioxidant system, consequently exacerbating neurodegeneration in PD. But not only microglia, even astroglia and maybe

Evidence of inflammation in Parkinson's disease	
Human data	
Increased levels of cytokines in the CSF and striatum in PD brains	Blum-Degen et al. 199
	Mogi et al. 1994a, b
	Muller et al. 1998
Increased number of activated microglia in PD brain	Banati et al. 1998
	Knott et al. 1999
	Imamura et al. 2003
	McGeer et al. 1988b
	Ouchi et al. 2005
	Sawada et al. 2008
Sustained microglial activity in humans exposed to MPTP years	Langston et al. 1999
Increased cytokine levels	Hunot et al. 1996
	Knott et al. 2000
	Mogi et al. 2000
	Mogi et al. 2007
	Reale et al. 2009
	Ros-Bernal et al. 2011
Parkinsonism models data	
Microglial activation in PD models	Barcia et al. 2004
	Carvey et al. 2005
	Cicchetti et al. 2002
	Gao et al. 2002
	Klintworth et al. 2009
	Greenamyre et al. 2003
LPS-induced microglial activation leads to dopaminergic	Chung et al. 2012
degeneration	Sui et al. 2009
	Zhang et al. 2010
	Zhang et al. 2012

Table 1 Most relevant articles correlating neuroinflammation and Parkinson's disease



Healthy dopaminergic neuron

**Fig. 4** Neuroinflammatory steps associated with dopaminergic cell death. Diverse initiation processes including gene mutations, brain injury, and alpha-synuclein deposits will activate immune cells in the CNS inducing microglial and astroglial activation, the production of cytokines (TNF- $\alpha$ , IFN- $\gamma$ , etc.), and the release of oxidative species such as ROS and RNS. This neuroinflammatory response may also include the involvement of components of the peripheral immune system and the alteration of the GC/GR pathway. As a result of this neuroinflammatory process, an environment of sustained increase of inflammatory and oxidative stress in the SNpc would be created. Alternatively, a healthy neuron may become dysfunctional as a result of alterations in the ubiquitin-proteasome system, direct exposition to neurotoxins, and the general oxidative stress inducing the dopaminergic cell death. Finally, this dying/dead dopaminergic neuron would secrete unknown signals that would induce phagocytosis by activated microglia (Adapted from Tansey et al. 2007)

oligodendrocytes have a main role in controlling neuroinflammatory processes in PD. In relation to the sustained microglia activation, GC-GR system is misbalanced in PD and contributes to the neuroinflammatory processes, where some genetic markers such as HLA-DR could be the new targets for drug development and pharmacogenetics against neurodegeneration (Fig. 4).

In sum, the inflammatory component of PD is having significant attention among researchers, and it will probably be assumed in the clinical scenario in the coming years. To date, there are numerous preclinical trials testing the protective effects of anti-inflammatory drugs in animal models of parkinsonism, and we expect that hopefully some of them will soon be brought into phase I trials. However, further researches on novel targets are needed to understand the specific aspects of neuroinflammation in PD. Acknowledgments This work was supported by grants from the Spanish Ministry of Science (FIS 2010–02827), Fundación Séneca (FS/05662/PI/07; FS/15329/PI/10), and CIBERNED (Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas). Finally, the authors declare that there is no conflict of interests in the present work.

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## **Neuromelanin and Parkinson's Disease**

## Giorgia Greco

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#### Abstract

Dark pigmented organelles, present in catecholaminergic neurons in specific brain regions, are indicated with the term neuromelanin (NM). They are complex structures, mainly comprised by granules of melanin polymer, closely associated with peptide and lipid components. Although NM is present in several animals, it is considered as a unique feature of the man due to the extremely higher degree of pigmentation, even in comparison to other primates. For a long time NM was considered an inert cellular waste product of poor interest that in the absence of mechanism of removal accumulates during the entire lifespan. Just recently, NM has received renewed attention for its role in Parkinson's disease (PD), where a

#### G. Greco

Analytical Research Group, Wissenschaftszentrum Weihenstephan, Technische Universität München, Freising, Germany e-mail: greco@wzw.tum.de

selective death of the NM-containing neurons of the substantia nigra (SN) pars compacta is observed, while nonpigmented neurons are mostly spared. A physiological accumulation of NM seems to be a protective phenomenon, which prevents several neurotoxic processes. In particular, in dopaminergic neurons of SN, where no ferritin has been detected, NM appears to function as an iron storage system. However, in PD patients NM released by dying neurons can trigger a vicious circle of neuroinflammation and ensuing neuronal death. This chapter presents the structure, the production, and the development of NM, as well as the recent hypotheses about physiological NM role and its behavior in pathological conditions.

### Keywords

5,6-dihydroxyindole • 5-S-cyteinyldopamine • Benzothiazine • Casing model • Catecholaminergic neurons • Dolichol • Dopaminequinone • Dopaminochrome • Eumelanin • Ferritin • Melanin, peripheral • Microglia activation • Mixed melanogenesis • Neuromelanin • NM, age • NM casing model • NM, degradation products • NM, electron density • NM, enzymatic control • NM, iron binding sites • NM, iron level • NM, iron storage system • NM, lipid component • NM membrane • NM, metal chelator • NM, neuron selective death • NM organelle • NM, organic molecule interaction • NM, oxidation potential • NM, peptide component • NM pigment • Pheomelanin • Pigmented neurons

#### List of Abbreviations 4-AHPEA 4-amino-3-hydroxyphenylethylamine AFM Atomic force microscopy DA Dopamine DAO Dopaminoquinone DOPA Dihydroxyphenylalanine DO DOPA-quinone FEL Free-electron laser NM Neuromelanin PD Parkinson's disease PDCA Pyrrole-2,3-dicarboxylic acid PEEM Photoelectron emission microscopy PTCA Pyrrole-2,3,5-tricarboxylic acid RNS Reactive nitrogen species ROS Reactive oxygen species SEM Scanning electron microscopy SN Substantia nigra TDCA Thiazole-4,5-dicarboxylic acid TTCA Thiazole-2,4,5-tricarboxylic acid

### 1 Introduction

Neuromelanin (NM) is a natural brown-black polymeric pigment present in the highest concentration in the catecholaminergic neurons of substantia nigra (SN) and locus coeruleus of human brain.

This pigment belongs to the melanin family, which members include the peripheral melanins, responsible in the mammalian for the coloration of hair, skin, and iris (Prota 1992).

NM has stimulated the interest of numerous scientists for almost one century. Pigmented neurons were observed in man since the 1930s, and it was long assumed that the intense pigmentation of selected brain regions was a unique feature of the human being and did not occur even in other primates (Adler 1939; Foley and Baxter 1958). Some years later, the pigment was observed in SN of the adult gorilla (Adler 1942), chimpanzee, orangutan, gibbon, certain old-world monkeys and lemurs (Scherer 1939), cat and dog (Brown 1943), and horse (Gillilan 1943). On the basis of these results, Marsden (1961) investigated systematically the distribution of the pigment in SN of mammals in 49 species from 11 mammalian orders, founding that the dark granules were present in the nigral cells of the members of the Primates, the Rodentia, the Carnivora, the Perissodactyla, the Artiodactyla, the Xenarthra, and the Marsupialia. The pigmented cells varied from animal to animal only in regard to their number and intensity of the pigmentation. Interestingly, the SN cells of all the examined 16 species of the Primate order were pigmented and more intensely respect to the members of the other orders, reaching the highest degree of pigmentation in man. However, the presence of the pigment in such a wide variety of mammalian orders did not lead to any indication to its function in the brain.

Therefore, for a long time NM was considered an inert cellular waste product of poor interest. Just in recent years, NM has received renewed attention for its role in Parkinson's disease (PD), one of the most common neurodegenerative disorders with clinical features among which bradykinesia, rigidity, tremor, and postural instability (Calne et al. 1992; Gelb et al. 1999). These classic motor symptoms result from the progressive and selective death of the NM-containing neurons (NM, neuron selective death) of the SN pars compacta, which form an integral part of the basal ganglia motor circuit (Double 2012). The loss of the NM-containing neurons of SN is the main pathological process observed in PD, while nonpigmented neurons are mostly spared (Bush et al. 2006), and this is considered a criterion required for a confirmed diagnosis of PD (Braak et al. 2003).

Many questions about the significance of NM are still awaiting answers, despite it may be of primary importance for providing information to the etiology of PD. Until now no unambiguous physiological role has been defined for NM, as well as the relationship between the presence of NM and the neuronal susceptibility has not yet been clarified. In addition, NM biosynthesis, whether it is enzymatically controlled or results from an autoxidation pathway, is still object of investigation.

In this chapter, the distribution, the structure, the biosynthesis, and the development of NM in the human brain are presented. Moreover, the recent hypotheses about NM physiological role, also in relation to its architecture, and NM pathogenic action in the selective degeneration of pigmented neurons during PD are discussed.

### 2 Distribution and Appearance of Neuromelanin (NM)

In man NM is mainly found in the cytoplasm of the catecholaminergic neurons of the SN pars compacta of the midbrain and in the neurons of the locus coeruleus within the pons, where almost all the neurons are pigmented and a darkened area can be seen macroscopically (Fedorow et al. 2005a). Some pigmented neurons are also found in other regions of the pons and the midbrain, in the hypothalamus, and medulla oblongata (Bazelon et al. 1967; Rosengren et al. 1985), where, however, the number of NM-containing cells is around tenfold lower than those in SN (Bogerts 1981).

Only dopamine (DA) and noradrenaline neurons are reported to contain NM, while no NM is preset in adrenaline neurons. Interestingly, within the brain not all the DA and noradrenaline neurons produce NM. The pigment is found in 95 % of the DA neurons of SN, only in 50 % of the neurons of the ventral tegmental area, and other DA neurons, as, for example, those of the olfactory bulb do not produce the pigment at all (Double et al. 2008). This suggests that even though DA or noradrenaline is required for NM biosynthesis, their presence within the neurons does not ensure NM production.

Due to the number of pigmented neurons in human SN pars compacta and the high concentration of pigment inside the cells, NM isolated from SN is the most studied pigment. It has been evaluated that NM levels in a middle-aged man without any neurological disorders are around 2  $\mu$ g/mg wet weight of SN (Zecca et al. 2002a).

Scanning electron microscopy (SEM) and atomic force microscopy (AFM) images of NM have revealed that the dark pigment is composed by granules with an average diameter of ~350 nm, comprised of small spherical particles of ~30 nm diameter (Bush et al. 2006). Similar morphological organization with substructures of spherical constituents with diameters on the order of 30 nm has been observed also for other members of the melanin family, among which the natural melanin from the ink sac of the cuttlefish *Sepia officinalis* and the melanin isolated from human hair and from bovine and human eyes (Liu and Simon 2003). Besides the natural occurring pigments, also SEM images of bioinspired synthetic melanin have revealed tightly aggregated globe-like nanosized particles with an average diameter of  $39 \pm 9$  nm, resembling the architecture of the natural melanins (Greco et al. 2011).

Differently from the other members of the melanin family, NM pigment is naturally found as organelles (NM organelle) comprising three different electrondense constituents (NM, electron density). The dark electron-dense component, identified as the melanin polymer itself, is closely associated with a component of intermediate electron density and an electron-lucent component, suggested to be lipid (Duffy and Tennyson 1965; Moses et al. 1966). The question of whether NM is surrounded by a membrane or not is still debated. Early studies clearly showed that the organelles are membrane bound (Moses et al. 1966; Hirosawa 1968), whereas other works did not confirm this observation (Duffy and Tennyson 1965). Recently, NM synthesis was induced in rat SN neurons and cell cultures by exposure to DOPA (dihydroxyphenylalanine). These model organelles were surrounded by a double membrane separated by  $\sim 15$  nm. NM isolated from human SN showed similar features, but no strong evidence for the existence of a double membrane was provided (Sulzer et al. 2000). The controversial information in several papers about NM membrane may be the consequence of different purification methodologies, which could alter the aspect of the granule. It has also been proposed that a lack of a clear evidence for a membrane enclosing the NM organelle may be due to the difficulties of preserving membranes in the human brain postmortem (Fedorow et al. 2005a). Further studies are necessary to clarify the presence of a membrane that may play an important role in moderating or even preventing the interaction of the pigment with other neuronal constituents.

A recent investigation has revealed the presence of dark pigmented areas in noncatecholaminergic neurons and in brain regions other than SN and locus coeruleus (Zecca et al. 2008). The organelles observed in neurons of the putamen, premotor cortex, and cerebellum exhibited three components of different electron density surrounded by a double membrane, resembling the NM of SN. The electron-dense component was comprised of substructures of 30 nm diameter and was identified as a member of the melanin family on the basis of its physical-chemical properties.

The pigment was termed NM too, even if it derives biosynthetically from the different precursor DOPA, instead of DA or noradrenaline. The study has shown that pigmented aggregates with common structure, behavior in aging, and interaction with metal ions and lipids, besides NM of SN and locus coeruleus, are ubiquitous within the human brain (Zecca et al. 2008). This result opens new and interesting questions about the function of melanin pigments in the human central nervous system. In particular, the peculiar NM architecture involving different constituents suggests that NM is not likely to be a waste product without significance, as considered until some years ago.

### 3 Structure of NM

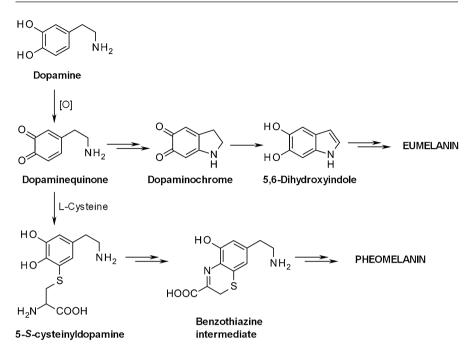
The present information about NM structure derives prevalently from studies performed by several independent research groups on NM isolated from human SN. In addition to the dark melanin pigment, peptide and lipid components concur to form the NM structures of different electron density observed in electron micrographs. The peptide component is covalently bound to the melanin structure and corresponds to ~15 % of the NM weight, while lipids are adsorbed on the surface of the pigment and account for ~20 % of the NM weight (Zecca et al. 2000). In the following paragraphs, the melanin pigment, the peptide, and the lipid components of NM granule are discussed in detail.

### 3.1 The Melanin Pigment

NM pigment is considered a member of the melanin family, as similarly to the pigment of skin, eyes, and hair, it is a heterogeneous low soluble aromatic polymer, with a stable free radical and high affinity for transition metals (Bridelli et al. 1999).

Several experimental evidences support the common view that in vivo NM derives from oxidation of DA (Double et al. 2008). Two complementary strategies, also used for the coetaneous melanin, have been employed to study the structure of the pigment. Naturally occurring NM has been investigated by different chemical degradation methods and following analysis of the liberated products, while the polymer formation has been mimicked in vitro by oxidation of DA and isolation of intermediate compounds (Wakamatsu et al. 2003; Tse et al. 1976; Young and Babbitt 1983; Zhang and Dryhurst 1994). At the present, in vitro oxidation of DA is commonly used for the preparation of synthetic pigments, as models of NM (Double et al. 2000; Zareba et al. 1995). Both degradative and biosynthetic strategies have shown that NM comprises two distinct classes of polymers: eumelanin and pheomelanin. The pathway for NM biosynthesis has been suggested to be similar to that for the synthesis of peripheral melanins (Ito and Wakamatsu 2006). The first step is DA oxidation to dopamine quinone (DAO) that for intramolecular cyclization and further oxidation is converted into dopaminochrome. This a reactive specie that can rearrange to 5,6-dihydroxyindole, which for polymeric oxidation leads to the eumelanin. Alternatively, DAO conjugates to L-cysteine, with ensuing formation of isomeric cisteinly dopamine adducts, mainly 5-S-cyteinyldopamine. This adduct for oxidation, followed by intramolecular cyclization of the cysteine chain, gives rise to benzothiazine units, which polymerize leading to the pheomelanin (Zhang and Dryhurst 1994) (Fig. 1).

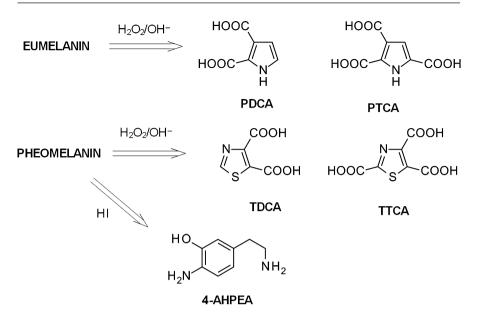
Peripheral eumelanin and pheomelanin pigmentation (Melanin, peripheral) has been more extensively studied. Both pigments derive from a common precursor, the DOPA-quinone (DQ). Eumelanin appears as a dark pigment responsible for the brown-black coloration of skin and hair, while pheomelanin determines the carroty to fire-red color of hair, commonly found in Celtic populations. Over the past 30 years, chemical analyses have been performed that enable the quantification of the amount of eumelanin and pheomelanin present in a naturally occurring sample of melanin. These works collectively reveal that more often than not, natural melanin pigments consist of both eumelanin and pheomelanin in varying ratios (Ito and Wakamatsu 2003; Thody et al. 1991). Recently, by the use of modern surface techniques, in particular photoelectron emission microscopy (PEEM) coupled to a free-electron laser (FEL), it has been shown that human black hair melanin has a surface oxidation potential of -0.2 V versus the normal hydrogen electrode, attributed to the eumelanin, while melanin form red hair has two surface oxidation potentials of -0.2 and +0.5 V, respectively. The negative potential has been ascribed to eumelanin pigment, whereas the positive to the pheomelanin one. This indicated that melanin from red hair has both types of melanin on or near their surface, and on the basis of the oxidation potential values, eumelanin is an antioxidant, while pheomelanin exhibits a prooxidant behavior (Ito 2006; Simon et al. 2009).



**Fig. 1** Formation of indole and benzothiazine units from DA oxidation and their polymerization to eumelanin and pheomelanin pigment of NM granule

Supported by biochemical evidence, a casing model (NM casing model) with a pheomelanin core encased in a eumelanin shell was proposed in 1982 and since then it has been sustained by different other studies (Agrup et al. 1982; Simon and Peles 2010). On the basis of this model and on the eumelanin/pheomelanin ratios, the melanin granule from red hair results to have a tinier eumelanic coat, exposing the pheomelanin core. In the last years epidemiological and clinical data support the hypothesis that eumelanin is photoprotective, defending the tissues from UV-induced damages, whereas pheomelanin may be the main cause of the abnormal susceptibility to erythema and skin cancer, observed in red hair people (Bliss et al. 1995).

Two microanalytical methods, initially developed to quantify the amount of eumelanin and pheomelanin in cutaneous tissues and hair, have been applied to NM isolated from SN (Wakamatsu et al. 2003). The first method consists in the alkaline hydrogen peroxide oxidation of the sample, followed by quantification of pyrrole-2,3-dicarboxylic acid (PDCA) and pyrrole-2,3,5-tricarboxylic acid (PTCA), degradation products of eumelanin (NM, degradation products), and thiazole-4,5-dicarboxylic acid (TDCA) and thiazole-2,4,5-tricarboxylic acid (TTCA), degradation products of pheomelanin. The second method is specific for pheomelanin and it is based on the quantification of 4-amino-3-hydroxyphenylethylamine (4-AHPEA), arising from reductive hydrolysis of the pigment with hydriodic acid (Fig. 2).



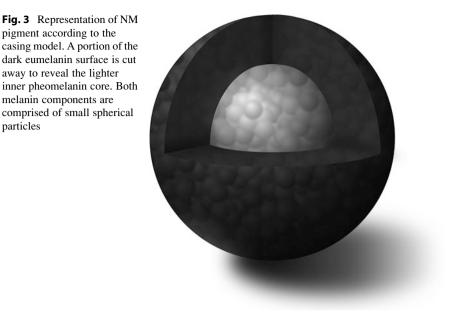
**Fig. 2** Degradation products of eumelanin and pheomelanin pigment by treatment of NM granule with alkaline hydrogen peroxide or hydriodic acid

These methods showed in NM isolated from SN the presence of eumelanin and pheomelanin in a ratio of 3:4 (Wakamatsu et al. 2003). FEL–PEEM analysis has revealed for NM granules a single oxidation potential (-0.1 V) (NM, oxidation potential), indicating the presence of only eumelanin on the surface; no superficial pheomelanin have been detected (Bush et al. 2006). Therefore, the eumelanin surface is built up on the pheomelanin core according to the casing model (Fig. 3).

X-ray diffraction studies have shown that NM granule of SN, as other natural peripheral melanin, has a multilayer three-dimensional structure, composed by planar overlapped sheets at about 4.7 Å of distance (Zecca et al. 2008). Just recently the pheomelanin core is receiving attention. By X-ray absorption spectroscopy, the atomic structure around the atom of S in natural and synthetic NM has been investigated, revealing for the first time in a direct and nondegradative way the presence in human NM of heterocyclic S of the benzothiazine type (Crippa et al. 2010).

### 3.2 Peptide Component in NM

Whereas several efforts have been directed to the comprehension of the pigment structure, only recently the presence of other components in NM granule has been investigated. Zecca et al. (2000) showed a peptide component in NM (NM, peptide component), after treatment with different strong agents for protein removal.



The peptides resulted to be covalently bound to the melanin structure, as shown by comparison with a treatment for eliminating only noncovalently bound proteins. The peptide residues were present even after digestion with proteolytic enzymes, indicating that the peptide chains are in some way sterically protected by the melanin pigment. Furthermore, synthetic melanin incubated in brain homogenate showed the ability to bind peptides chemically.

The cysteine units in the intact bound peptide chain represent  $\sim 3 \%$  of the entire amino acid content, and their amount is almost unchanged in the strongly bound peptide residues, suggesting that they are localized in proximity of the pigment. The association of melanin to peptide component may result from a direct reaction of catechol moieties of the indole units on the surface of NM granule with the cysteine residues of peptides. At support to this interpretation, a model study has shown the formation of protein-bound cysteinylcatechols by incubation of DA or norepinephrine with bovine serum albumin (Ito et al. 1988). Proteomics of NM granules (NM, proteomic) has allowed the identification of more than 70 proteins, occurring commonly in human brain tissues, mainly lysosomal proteins, sustaining the hypothesis of a genetic program for the formation of NM granules (Tribl et al. 2005).

### 3.3 Lipid Component in NM

The electron-lucent component comprises 35 % of the NM granule volume (Fedorow et al. 2006) and contains neutral lipids (NM, lipid component). A twostep washing of the granules with methanol and hexane has allowed a completed removal of the lipid component, proving that they are adsorbed onto the surface of the granule and not chemically bound (Zecca et al. 2000). Interestingly the major lipid component of NM is the polyisoprenoid dolichol, which contains 17–23 isoprenoid units and accounts for approximately 12 % of NM weight (Fedorow et al. 2005b; Ward et al. 2007). Other lipids have been also found, as cholesterol, ubiquinone-10,  $\alpha$ -tocopherol, and dolichyl esters, but they are present only in small quantity (Fedorow et al. 2005b).

Small amounts of dolichol are ubiquitously distributed in the body, generally located in cellular membranes, where it is supposed to have a role in the stabilization of the membrane itself. While the phosphorylated form of dolichol is involved in the formation of glycoproteins in the endoplasmatic reticulum (Varki et al. 1999), the function of free dolichol remains to be clarified.

Accumulation of dolichol is associated with neurodegeneration in disorders such as the neuronal ceroid lipofuscinoses (Kin et al. 1983). In NM granule, the presence of large dolichol deposits represents until now the first example of dolichol concentrated in intracellular inclusions in healthy tissue (Fedorow et al. 2006). The association of dolichol with NM results to be specific. Indeed cholesterol, which in all the other brain regions is typically 200-fold higher than dolichol on a mass basis (Andersson et al. 1987), is present at considerably lower concentration in NM lipid. Dolichol may play a structural function, as it can alter membrane lamellar bilayer into nonlamellar or hexagonal structure (Zhou and Troy 2004).

Recently, the influence of NM lipids in the aggregation of the granule has been investigated (Dedov et al. 2007). The removal of all NM lipids resulted in significantly larger granules than those present prior to lipid removal. Interestingly, selective removal of cholesterol by methanol extraction resulted in an eight- to tenfold enhancement in granule size, whereas the pro-aggregation effect of dolichol extraction using hexane was significantly smaller. These data showed that NM lipids, preventing the formation of large pigment aggregates, may be a factor regulating NM granule size in vivo (Dedov et al. 2007).

Under oxidative stress condition, lipid peroxidation appears to contribute to neurodegeneration in PD (Dexter et al. 1989). In this contest, dolichol may be a substrate of the lipid peroxidation, as in vitro studies have shown a time-dependent degradation of dolichol when incubated with a radical initiator (Dedov et al. 2007). These observations are just preliminary and further investigations are required for clarifying the structural and physiological role of NM lipids in normal and pathological conditions.

### 4 Biosynthesis of NM

The study of neuromelanogenesis is made more complicated than that of peripheral melanin biosynthesis, because NM is absent from all of the commonly used laboratory animals including rodents. A general understanding of the first steps of NM biosynthesis has been provided by investigation of the products arising from in vitro biomimetic oxidation of NM precursors and comparison with what is

known about peripheral melanins (Fig. 1). It is extensively debated whether the NM synthesis is enzymatically controlled (NM, enzymatic control), like all melanins in the periphery, or whether NM arises from a simple autoxidation of DA. Although DA cells in the midbrain express tyrosine hydroxylase, the rate-limiting enzyme for DA synthesis, there is no correlation between the degree of pigmentation and tyrosine hydroxylase immunoreactivity (Gaspar et al. 1983). The enzyme tyrosinase is a key enzyme in the synthesis of peripheral melanin, since it catalyzes the conversion of the amino acid tyrosine to DQ, the precursor of eu- and pheomelanin (Korner and Pawelek 1982). However, immunochemistry studies have no detected tyrosinase in human midbrain (Ikemoto et al. 1998). At support to that, albinos, who are characterized by absence of peripheral pigmentation due to a lack of functional tyrosinase, have normally pigmented neurons in SN (Foley and Baxter 1958). Also the enzyme peroxidase and macrophage migration inhibitory factor have been proposed for NM synthesis. However, their distribution in the brain does not correlate to NM distribution (Okun 1997; Matsunaga et al. 1999). Several other enzymes have been suggested to be associated with NM biosynthesis, but to date no likely candidate has been demonstrated to be involved in the neuromelanogenesis. In the absence of a convincing enzymatic route, for a long period NM has been regarded as a waste product of catecholamine metabolism (Bogerts 1981; Graham 1979; Mann and Yates 1983; Mann et al. 1977). Indeed, NM is not present at birth and increases over the lifespan, supporting the idea that the pigment is a cellular by-product of DA metabolism formed via autoxidation and, in the absence of any mechanism to remove it, accumulates intracellularly with age. However, several evidences are in contrast with this view. First of all, NM and DA do not have the same cellular distribution in human brain and the pigment seems to be a peculiar feature of the man, lacking or at least present in small amount in the midbrain of many other species (Fedorow et al. 2005a). In addition, PD patients treated with DOPA, which it is rapidly converted in DA by the enzyme aromatic acid decarboxylase, do not exhibit increased quantities of NM within their surviving SN neurons as might be expected to be the case if NM represents a product of pure DA autoxidation (Zecca et al. 2001a). Finally, the complex casing structure proposed for the NM pigment and its aggregation with specific protein and lipid components support the existence of a controlled process.

The casing model may be the consequence of a mixed melanogenesis regulated in some way by cysteine levels. In vitro studies on biosynthesis of peripheral melanin have clearly indicated that DQ conjugates to cysteine leading to pheomelanin production until cysteine concentration is above 1  $\mu$ M. At lower cysteine levels DQ polymerizes in eumelanin (Ito 2003; Land et al. 2003). This favored production of pheomelanin over eumelanin suggests that the first step is the formation of the pheomelanin core, and then, when the cysteine is depleted, eumelanin production starts. In NM production, this casing process is further favored by the fact that DAQ cyclizes much more slowly than DQ, leading to a preferential production of cysteinyldopamines (Ito 2003). The construction of NM granule seems to be a regulated process in such a manner that DAQ is conjugated first with cysteine to form cysteinyldopamine isomers, which are further oxidized to form a pheomelanic core. This step is followed by the production of a eumelanic surface (Ito 2006). Whereas it is not well understood how initially DA and later the cysteinyldopamines are oxidized for producing the pheomelanin pigment, the synthesis and the following deposition of eumelanin pigment onto the preformed pheomelanic core may be a spontaneous process, in part catalyzed by the prooxidant pheomelanin core, as observed in biomimetic studies (Greco et al. 2011).

### 5 Developmental Stages of NM

NM concentration in SN pars compacta increases similarly in men and women over the entire lifespan. In the first year of life, NM is in general not detectable. Between 10 and 20 years, the NM levels are  $0.3-0.8 \ \mu\text{g/mg}$  of SN, between 20 and 50 years are  $0.8-2.3 \ \mu\text{g/mg}$  SN, and between 50 and 90 are  $2.3-3.7 \ \mu\text{g/mg}$  of SN (Zecca et al. 2002a) (NM, age).

In accordance with these quantitative analyses, light microscopy images have revealed that no NM is visible in SN neurons in the prenatal or infant brain. The pigment first appears within the cytoplasm of the DA neurons at around the age of 3 years as small granules, whose size and number increase until the age of 20. At this time NM fills, on average, 47 % of the cytoplasm. After the age of 20, the proportion of the cell occupied by the pigment remains stable, but the amount of pigment increases with age (Halliday et al. 2006). Over time the coloration and the appearance of NM change too. Initially, NM occurs as dusty yellow-brown granules, which progressively form dark denser clusters (Halliday et al. 2006). These data may represent a direct observation of the construction of the NM granule in vivo according to the casing model, where the fine yellow-brown pheomelanin is form first and then it is encased by the dark eumelanin shell. However, at the present no evidence of the pigment in the on-growing granule is available. In this contest, further works, leading to clarify the variation of eumelanin/pheomelanin levels with aging, will be a fundamental step in the study of neuromelanogenesis.

Interestingly, also brain dolichol levels increase over the human lifespan. At the age of 6 years, dolichol in human brain gray matter is around 18  $\mu$ g/g wet weight, exceeding the 200  $\mu$ g/g after the sixth decade of life (Pullarkat and Reha 1982).

### 6 Dual Role of NM

In contrast to peripheral melanin, whose main function in the body is to protect underlying tissues from harmful UV radiation (Ito 2006), much less is known about the role of NM.

No physiological function has been defined for NM and two different roles have been suggested, either protective or toxic with respect to the degeneration of catecholaminergic neurons depending on the cellular context. An increasing number of new studies concord with the view that NM is a cellular protecting system in the normal brain. NM synthesis protects the neurons in the SN, because the melanin component is generated from cytosolic catechols which are not accumulated in synaptic vesicles (Zucca et al. 2004). Therefore, this mechanism ensures the removal of reactive/toxic quinones arising from DA oxidation that otherwise would cause neurotoxicity (Paris et al. 2010). Furthermore, NM can trap a variety of organic molecules, as drugs (e.g., chlorpromazine, haloperidol, imipramine), pesticides (e.g., paraquat), and toxic compounds (e.g., 1-methyl-4-phenylpyridinium (MPP<sup>+</sup>), beta-carbolines), and immobilize them as stable adducts, thus protecting against toxicity (Salazar et al. 1978; Lindquist et al. 1988; Ostergren et al. 2004; D'Amato et al. 1986) (NM, organic molecule interaction).

Similar to peripheral melanin, NM is a strong chelator of metal cations such as iron, zinc, copper, manganese, chromium, cobalt, mercury, lead, and cadmium (Liu et al. 2004; Zecca and Swartz 1993; Zecca et al. 2002b) (NM, metal chelator). In NM isolated from SN, iron is the metal occurring at the highest concentration, with levels above 10  $\mu$ g/mg NM (NM, iron level). In DA neurons of SN ferritin are not present, whereas it has been found in glial cells (Zecca et al. 2004; Moos 2000). Therefore, nowadays NM is the only well-documented storage system of iron in SN neurons (Zucca et al. 2004) (NM, iron storage system). In normal condition, NM can sequester iron ions, playing an important role in the protection the neurons from iron-induced oxidative stress (Zareba et al. 1995). NM contains both high- and lowaffinity iron-binding sites, where the higher-affinity binding sites (NM, ironbinding sites) are particularly better able to sequester iron (Double et al. 2003). The pigment in vivo is suggested to be only partially saturated with iron, thus maintaining a residual chelating capacity to protect SN against iron toxicity (Shima et al. 1997). In the presence of high iron levels, NM accumulates iron in low-affinity binding sites, arranged in a ferritin-like oxyhydroxy iron clusters, where it can be redox active, producing free radicals and thus leading to neurotoxic consequences (Gerlach et al. 1995; Double et al. 2003; Zecca et al. 2001b). In parkinsonian subjects, where an abnormal loss of NM-containing neurons is observed, the NM levels are  $1.2-1.5 \,\mu$ g/mg of SN pars compacta, which is less than 50 % with respect to the age-matched controls (Zecca et al. 2002a). At support to NM protective role, the levels of redox activity are significantly higher in PD patients (+69 %) and increase in patients with the most severe neuronal loss (Faucheux et al. 2003).

On the other hand, NM can be also toxic in PD patients when it is released by dying neurons of SN. Insoluble extracellular NM is phagocytosed by microglia with consequent induction of neurotoxicity by overactivation of microglia (microglia activation) and ensuing production of reactive oxygen species (ROS) and nitrogen species (RNS), and proinflammatory factors (Zhang et al. 2011). Recent epidemiological studies have suggested that neuroinflammation, in particular in a chronic state, is recognized to contribute to the pathogenesis of PD (Chen et al. 2003; Gao et al. 2011; Wu et al. 2002). An initial damage of pigmented neurons, independently if the factors are genetic or environmental, causes the release of NM, which could trigger a vicious circle of neuronal death that enhances the inflammation. Moreover, as ROS can degrade NM, the consequence may be that the neurons not only lose

a protective agent but also release iron and other cytotoxic metals and compounds, accelerating the neuronal death (Zucca et al. 2004; Shima et al. 1997).

In addition to these studies, there are strong evidences for an alteration of NM structure since the first stages of PD. In particular, it has been observed during the disease a decrease of the lipid component and an increase of the density of the pigment (Halliday et al. 2005). Investigations by magnetic spectroscopies of the pigment structure have revealed that the melanin component undergoes modification (Fasano et al. 2006). Other studies have also showed that  $\alpha$ -synuclein protein, a primary component of Lewy bodies, aggregates on NM in SN in parkinsonian, but not in healthy control, brains (Halliday et al. 2005). Therefore, this interaction may explain the higher susceptibility during the disease process of highly pigmented brain regions, such as SN (Double 2012).

### 7 Implication of NM Architecture in Parkinson's Disease

The casing model reported for NM pigment is particularly intriguing and growing efforts are directed to the study of its formation and function in the brain.

In healthy conditions, the pheomelanin core of NM pigment may be the result of a detoxification process of DAQ through conjugation with cysteine. In fact, SN is the brain region with the highest levels of cysteinyldopamine isomers, the pheomelanin precursors (Spencer et al. 1998). The pheomelanin core may function as scaffold for the eumelanin pigment. Moreover, recent evidences suggest that it could also catalyze eumelanin formation (Greco et al. 2011). The eumelanin surface of NM granule is responsible for neuronal protection. Indeed, it is much more efficient than pheomelanin in binding drugs and metals ions (Mårs and Larsson 1999). In addition, the surface oxidation potential found for the eumelanin is not sufficiently to generate a high level of oxidative stress (Bush et al. 2006). On the other hand, any damage of the protective eumelanin surface by ROS, process that likely occurs in the human brain during PD, could expose the pheomelanin core. Pheomelanin, differently from eumelanin, has well-documented prooxidant property (Ye et al. 2006; Bush et al. 2006); therefore, the exposure of pheomelanin core may induce of a cascade of neurodegenerative events (Ito 2006).

At support to the fact that NM architecture may be involved in some way in the pathogenesis of PD, Gao et al. (2009) have recently reported an interesting correlation between the risk of PD and the hair color. They have shown that individuals with red hair have approximately a twofold higher risk for PD relative to those with black hair. It is known that the peripheral pigment granule of red hair individuals is characterized by a tiny eumelanin coat with consequent permanent exposure of the prooxidant pheomelanin core and this seems to be related to the higher incidence of melanoma in comparison to individuals with black hair (Samokhvalov et al. 2005; Vincensi et al. 1998). These findings should stimulate further researches directed to study of the NM architecture and the changing of melanin composition, in particular the eumelanin/pheomelanin ratio, with age and in pathological conditions.

### 8 Conclusion

Despites the recent progresses in the comprehension of NM biology, there are still many open questions. The future challenges are the clarification of the NM biosynthesis, its mechanism of turnover, and the role in heath individuals. Because of the complexity of NM granules, major insights into the chemical structure of the pigment, the peptide, and the lipid components would elucidate the possible interactions with other neuronal constituents. Moreover, a detailed understanding of NM structural and physiological changes in pathological conditions, in particular the contribution of NM to the oxidative stress and to neuronal vulnerability, would advance our knowledge of the underlying mechanisms that govern the progression of PD.

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# Neurotoxicity of 1-methyl-4-phenyl-1,2,3,6tetrahydropyridine (MPTP) as a Parkinson's Disease Model

H. Nakayama, T. Ito, Y. Shibui, T. Sai, K. Uchida, and X. J. He

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### Abstract

Administration of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) to susceptible animals is the most popular means for modeling the destruction of the nigrostriatal dopaminergic neurons seen in Parkinson's disease (PD). MPTP causes a damage to the substantia nigra pars compacta (SNpc) dopaminergic neurons and depletes striatal dopamine secretion in nonhuman primates and rodents. MPTP passes through the blood–brain barrier (BBB) and is converted, mainly in glial cells, into its active form, 1-methyl-4-phenylpyridinium (MPP<sup>+</sup>), by an enzyme, monoamine oxidase B (MAO-B). MPP<sup>+</sup> is selectively transported into dopaminergic nerve terminals through dopamine transporter (DAT) and finally induces dopaminergic cell loss. The selective neurotoxicity of MPTP to

H. Nakayama (🖂) • T. Ito • Y. Shibui • T. Sai • K. Uchida • X.J. He

Department of Veterinary Pathology, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan

e-mail: anakaya@mail.ecc.u-tokyo.ac.jp; t-ito@nirs.go.jp; maron2005@live.jp; Takafumi\_Sai@nts.toray.co.jp; auchidak@mail.ecc.u-tokyo.ac.jp; aaa31172@yahoo.co.jp

neural tissues other than the dopaminergic system, such as the subventricular zone (SVZ) and rostral migratory stream (RMS) in the adult brain, has been indicated. The cells undergoing apoptosis in the SVZ are A cells and MAO-B inhibitors completely protected against the neurotoxicity, suggesting that the MPTP neurotoxicity to A cells is mediated by the conversion of MPTP into MPP<sup>+</sup> by MAO-B. The neurotoxicity of MPTP to neuroblasts in the SVZ does not require DAT or other monoamine transporters. There is a strain difference in the susceptibility of the MPTP neurotoxicity between susceptible C57BL/6 and resistant BALB/c mice. However, MAO-B, DAT, neural NOS (nNOS), and inducible NOS (iNOS) expression levels do not influence the strain susceptibility to MPTP. MPTP and MPP<sup>+</sup> can pass through the placenta and embryonic or newborn BBB, and finally reach the brain. The number of MAO-B-positive glial cells in the brain increases in MPTP-treated embryonic and newborn mice, indicating the involvement of MAO-B in the acute neurotoxicity. MPTP or MPP<sup>+</sup> administration into embryonic and newborn mice causes a loss of TH-positive cells and fibers in the nigrostriatal system and increases the number of apoptotic cells in the SVZ. Delayed regeneration of dopaminergic neurons due to the loss of SVZ A cells may be one of possible causes of the pathological condition. With increased attention to dopaminergic neurogenesis in MPTP-induced animal models, such animals become useful as a tool for establishing the contact to PD on neurogenesis.

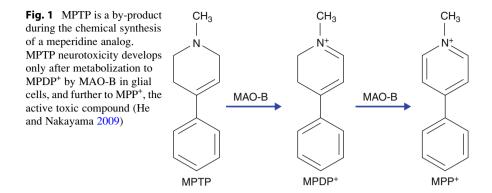
List of Abbreviations			
6-OHDA	6-Hydroxydopamine		
BBB	Blood-brain barrier		
BrdU	Bromodeoxyuridine		
CNS	Central nervous system		
DA	Dopamine		
DAT	Dopamine transporter		
DCX	Doublecortin		
DG	Dentate gyrus		
icv	Intracerebroventricularly		
iNOS	Inducible NOS		
MAO-B	Monoamine oxidase B		
MMPs	Metalloproteinases		
MPDP <sup>+</sup>	1-Methyl-4-phenyl-2,3-dihydropyridinium		
$MPP^+$	1-Methyl-4-phenylpyridinium		
MPTP	1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine		
nNOS	Neural NOS		
NOS	Nitric oxide synthase		
OB	Olfactory bulb		
PCNA	Proliferating cell nuclear antigen		
PD	Parkinson's disease		
RMS	Rostral migratory stream		
Ro 16-6491	N-(2-aminoethyl)-4-chlorobenzamide		
SERT	Serotonin transporter		

SGZ	Subgranular zone
SN	Substantia nigra
SNpc	Substantia nigra pars compacta
SVZ	Subventricular zone
TH	Tyrosine hydroxylase
TUNEL	Terminal deoxynucleotidyl transferase-mediated dUTP nick end
	labeling
VMAT2	Vesicular monoamine transporter 2

### 1 Introduction

Parkinson's disease (PD) is an age-related degenerative disorder of the central nervous system (CNS). It is often characterized by muscle rigidity, tremor, a slowing of physical movement (bradykinesia), and, in extreme cases, a loss of physical movement (akinesia). The symptoms of PD are attributed to the loss of pigmented dopamine-secreting (dopaminergic) neurons in the pars compacta region of the substantia nigra (SNpc) and a subsequent striatal deficiency of dopamine (DA). The idiopathic PD is also pathologically characterized by the presence of cytoplasmic neuronal inclusions, called Lewy bodies, in the affected region of the brain. The etiology of the disease has been poorly understood, and novel non-dopaminergic therapeutic strategies still remain challenging since the initial description was first described by James Parkinson in 1817 (Parkinson 1817). Although pharmacological dopamine replacement strategies provide a temporary symptomatic relief, there are at present no therapeutic methods for stopping the progressive neuronal cell loss (Borta and Hoglinger 2007).

Animal models are important tools in elucidating mechanisms involved in the pathological process and in investigating the new therapeutic strategies in PD. Currently, both genetic and toxic models of PD are available, but the use of neurotoxins such as 6-hydroxydopamine (6-OHDA), paraquat, 1-methyl-4-phenyl-1,2,3, 6-tetrahydropyridine (MPTP), and rotenone are still the most popular means for modeling the destruction of the nigrostriatal dopaminergic neurons seen in PD (Przedborski and Ischiropoulos 2005). Among them, 6-OHDA and MPTP have been more extensively used by investigators to produce PD models despite their limitations. 6-OHDA is a structural analog of catecholamines, DA and noradrenaline, and exerts its toxic effects on catecholaminergic neurons (Simola et al. 2007). Since 6-OHDA does not penetrate the blood-brain barrier (BBB) in adult rats, it must be administrated stereotactically into the substantia nigra (SN) or striatum to damage the dopaminergic nigrostriatal system (Perese et al. 1989; Przedbroski et al. 1995). Recognition of MPTP as a neurotoxin occurred early in 1982, when several young drug addicts mysteriously developed a profound Parkinsonian syndrome after the intravenous use of street preparations of meperidine analogs that, unknown to anyone, were contaminated with MPTP (Przedborski and Vila 2003; Langston et al. 1983), an incidental by-product during the chemical synthesis of a meperidine analog (Fig. 1)



with potent heroin-like effects (Przedborski and Vila 2003). MPTP has been found to produce irreversible Parkinsonism in humans almost indistinguishable from PD (Langston et al. 1983; Blume 1983). It is also well known that MPTP depletes striatal dopamine and causes a damage to the SNpc dopaminergic neurons in nonhuman primates and several species of rodents (Burns et al. 1983; Langston et al. 1984; Heikkila et al. 1984; Chiueh et al. 1984).

Recently, a great deal of interest has been focused on the stem cell therapies for PD. In addition to the fetal nigral transplantation, the capability of self-repairing in central nervous system (CNS) in adult mammalian has also been considered as a new insight into the cell-based approach for the treatment of neurodegenerative disorders. In PD or its animal models, it has attracted much attention whether the depletion of dopaminergic neurons triggers an activation of neural stem cells and their subsequent migration into the damaged area, and finally results in repopulation of dopaminergic neurons in SN. We have recently reported several results (He et al. 2006, 2008a, b; He and Nakayama 2009; Popolo et al. 2004), which will provide evidences available for endogenous neurogenesis in response to neurotoxin damages, and provide perspective on the contact of neurotoxin-based animal models on PD. This chapter will mainly describe such MPTP-induced neural damages in the subventricular zone (SVZ) as well as SNpc and striatum, and subsequent neurogenesis in the brain of the mouse. In addition, the strain difference in the susceptibility to MPTP in the mouse, and the relations of MPTP neurotoxicity and aging are mentioned.

### 2 Neurotoxicity of MPTP to Dopamine Neurons

MPTP is a by-product of the chemical synthesis of a meperidine analog, with potent heroin-like effects (Przedborski and Vila 2003), that causes damage to dopaminergic neurons and depletes dopamine in a manner similar to that seen in PD. When administered to animals, MPTP passes through the BBB and is converted, mainly in glial cells, into its active form, 1-methyl-4-phenylpyridinium (MPP<sup>+</sup>), by an enzyme, monoamine oxidase B (MAO-B) (Blum et al. 2001). For the last two decades, MPTP has been widely

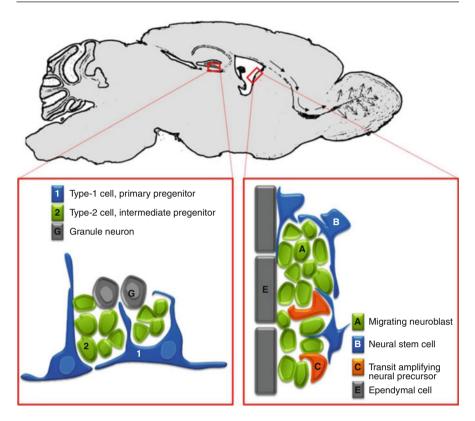
used to generate animal models of PD in rodents and nonhuman primates. A few studies have confirmed that MPTP can induce apoptosis of dopaminergic neurons in vivo (Tatton and Kish 1997; Vila et al. 2001; Spooren et al. 1998). When injected intraperitoneally (ip) with a subacute regimen, which consists of one injection per day (30 mg/kg per day) for 5 consecutive days, MPTP induces dopaminergic cell loss by a mechanism mainly involving apoptosis (Przedborski and Vila 2001). The use of this subacute regimen has been the main approach to studying the apoptosis of dopaminergic cells in mice. The induction of apoptosis in mouse nigrostriatal glia has also been demonstrated with this regimen (Serra et al. 2002), indicating that neurotoxicity of MPTP is not specifically restricted to dopaminergic neurons in the mouse brain. Furthermore, results from studies in vitro show that MPTP or MPP<sup>+</sup> induces apoptosis in primary cultured cerebellar granular neurons (Dipasquale et al. 1991; Du et al. 1997), mesencephalic dopaminergic neurons (Mochizuki et al. 1994), and neuroblastoma cells (Itano and Nomura 1995; Sheehan et al. 1997) as well as PC12 cells (Hartley et al. 1994; Desole et al. 1997).

### 3 Neurotoxicity of MPTP to Neural Stem Cells

### 3.1 The Origin and Properties of Neural Stem Cells

Neurogenesis continues into adult life in the brain of rodents (Altman 1963; Privat and Leblond 1972), nonhuman primates (Gould et al. 1999a, b; Pencea et al. 2001), and humans (Bernier et al. 2000; Eriksson et al. 1998) (Fig. 2). Neurogenesis has been confined largely to two discrete areas, the subventricular zone (SVZ), and the subgranular zone (SGZ) of the dentate gyrus (DG) (Garcia-Verdugo et al. 1998).

The SVZ, located throughout the inner wall of the lateral ventricle, is the largest germinal region and harbors neural stem cells that retain the capacity to generate multiple cell types (Doetsch et al. 2002). The SVZ is composed of neural progenitor cells (migrating neuroblasts, A cells), neural stem cells (astrocytes, B cells), neural precursor cells (rapidly dividing transit-amplifying cells, C cells), and ependymal cells (Doetsch et al. 1997, 1999). Neural stem B cells divide to give rise to clusters of precursor C cells, which in turn generate neuroblast A cells (Doetsch et al. 1999). Newly generated A cells in the SVZ migrate through a network of tangential pathways in the lateral wall of the lateral ventricle and then converge onto the rostral migratory stream (RMS) to enter the olfactory bulb (OB) with a behavior of "chain migration" (Doetsch and Alvarez-Buylla 1996), where they differentiate into granule cells and interneurons (Doetsch and Alvarez-Buylla 1996; Lois and Alvarez-Buylla 1994). It has been suggested that prokinetic n 2 serves as a chemoattractant for SVZ-derived neuroblast A cells, which appears to guide the migration of A cells from the SVZ through the RMS to their final layers in the OB (Ng et al. 2005). The multipotential A cells reside in the SVZ and RMS are eliminated through apoptosis to maintain a balance for proper development of the mammalian nervous system (Brunjes and Armstrong 1996; Levison et al. 2000;



**Fig. 2** Schematic representation showing the migratory pattern of newly generated A cells in the SVZ (*up*), and the cell population in the SGZ of DG (*bottom left*) and SVZ (*bottom right*) in adult rodent brain (He and Nakayama 2009)

Biebl et al. 2000). Metalloproteinases (MMPs) may also play a crucial role in the migration of individual A cells since the migration rates are reduced by the presence of inhibitors of MMPs (Bovetti et al. 2007). The architecture and function of the adult human SVZ differs significantly from that described in other mammals. There are four layers with varying thickness and cell densities throughout the lateral ventricular wall: a monolayer of ependymal cells (Layer I), a hypocellular gap (Layer II), a ribbon of cells (Layer III) composed of astrocytes, and a transitional zone (Layer IV) into the brain parenchyma (Quinones-Hinojosa et al. 2006).

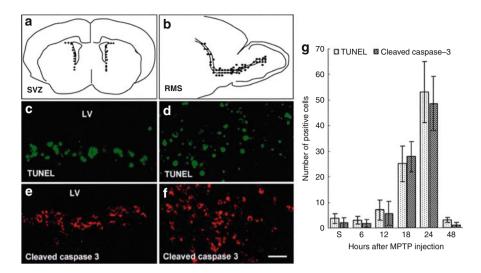
The DG is a part of the hippocampal structure. Neuronal progenitors in the adult mammalian hippocampal DG reside in the SGZ, which is located in the hilus immediately beneath the granular layer of the DG (Kaplan and Hinds 1977; Cameron et al. 1993). Neurogenesis in the DG was first demonstrated 40 years ago by autoradiography in rodents (Altman and Das 1965), and thereafter was further demonstrated in all mammalian species including human (Eriksson et al. 1998) and nonhuman primates (Gould et al. 1998, 1999a, b). Two types of neural progenitors can be identified in the SGZ according to their specific morphologies

and expression of unique sets of molecular markers (Zhao et al. 2008). The primary progenitors (type-1 cells) have the appearance of radial glia, which also express glial fibrillary acidic protein (GFAP). They share the similar features with type B cells residing in the adult SVZ and then are suggested as the putative stem cell population (Seri et al. 2001). Type-2 cells (intermediate progenitors) are GFAP negative and highly proliferative (Kempermann et al. 2004), sharing similar features with neuroblast A cells residing in the adult SVZ. Type-2 cells may arise from type 1-cells, but direct evidence delineating this lineage relationship is still lacking (Zhao et al. 2008). The newborn type-2 cells disperse and migrate a short distance into the granule cell layer where they differentiate, extend axons, and express neuronal marker proteins (Cameron et al. 1993; Kuhn et al. 1996). The migration of newborn neurons in the DG may also be controlled by guidance cues, as these cells only migrate to the hilus or the molecular layer under pathological conditions, such as in animal models of temporal lobe epilepsy (Zhao et al. 2008).

### 3.2 Neurotoxicity of MPTP to SVZ Neural Stem Cells

MPTP neurotoxicity develops only after metabolization to 1-methyl-4-phenyl-2,3-dihydropyridinium (MPDP<sup>+</sup>) and further to 1-methyl-4-phenylpyridinium (MPP<sup>+</sup>), the active toxic compound, by MAO-B (Fig. 1) (Przedborski and Vila 2003). Since MPP<sup>+</sup> is selectively transported into presynaptic dopaminergic nerve terminals through the dopamine transporter (DAT) (Chiba et al. 1985; Javitch et al. 1985; Gainetdinov et al. 1997; Bezard et al. 1999; Pifl et al. 1993), and the absence of the DAT on dopaminergic neurons confers a complete protection against MPTP toxicity (Gainetdinov et al. 1997; Bezard et al. 1999), the neurotoxicity of MPTP is suggested to be selective to dopaminergic neurons in SN. MPTP is mainly used in nonhuman primates and in rodents (Gerlach and Riederer 1996; Przedborski et al. 2001). In rodents, the C57BL/6 mice are widely used to create the PD animal model for understanding the disease pathogenesis in a great deal of reports despite their limitations.

In recent years, the selective neurotoxicity of MPTP to neural tissues other than the dopaminergic system in the adult brain has been elucidated. Several studies (Serra et al. 2002; He et al. 2006, 2008a) have demonstrated that MPTP also destroys forebrain migrating neuroblasts and nigrostriatal glial cells (astrocytes) in the adult mouse brain. In our previous studies (He et al. 2006, 2008a), the number of apoptotic cells in the SVZ and RMS peaked at 24 h after MPTP-injections, and decreased thereafter, paralleling the change in the number of cleaved caspase-3-positive cells (Fig. 3). The cells undergoing apoptosis in the SVZ, RMS, and OB were identified as A cells using immunohistochemistry and ultrastructural analyses, while a few were astrocytes (B cells) and none were transitamplifying precursors (C cells) (He et al. 2006, 2008a). The decrease in A cell numbers was most marked at day 2 and lasted to day 8 after the administration (He et al. 2006, 2008a). We also demonstrated that MAO-B inhibitors, such as deprenyl or N-(2-aminoethyl)-4-chlorobenzamide (Ro 16-6491), completely protected



**Fig. 3** Detection of apoptotic cells in the SVZ (coronal section) and RMS (sagittal section) with fluorescence TUNEL and immunofluorescence staining for cleaved caspase-3. (**a**, **b**) Schematic distribution of apoptotic cells in the SVZ and RMS with the positions of the cells undergoing apoptosis indicated by *dots*. (**c**, **d**) Apoptotic nucleus (*green*) in the SVZ and RMS revealed by the TUNEL technique, and (**e**, **f**) cleaved caspase-3 staining (*red*) in the same field. Scale bar =  $50 \mu m$ . (**g**) The results of a quantitative analysis of TUNEL-positive or cleaved caspase-3-immunoreactive cells in the SVZ of MPTP-treated mice. The number of TUNEL-positive or cleave caspase-3-immunoreactive cells peaked at 24 h and had declined to very low levels by 48 h after MPTP injection. Neither TUNEL-positive nor cleaved caspase-3-immunoreactive cells were found in control animals at any time point. Abbreviation: *LV* lateral ventricle (He et al. 2006)

against MPTP neurotoxicity to A cells, suggesting that MPTP-neurotoxicity to A cells is also mediated by the conversion of MPTP into MPP<sup>+</sup> by MAO-B (He et al. 2008b).

### 3.3 Neurotoxicity of MPTP to Neural Stem Cells Is Not Regulated by Monoamine Transporters

In dopaminergic neurons, MPP<sup>+</sup> enters into the cytoplasm via the dopamine transporter (DAT) (Gainetdinov et al. 1997; Javitch et al. 1985), is concentrated by an active process within the mitochondria, and impairs mitochondrial respiration by inhibiting complex I of the electron transport chain (Vila and Przedborski 2003). DAT is thus thought to play a crucial role in the MPTP neurotoxic process.

On the other hand, other monoamine transporters, including vesicular monoamine transporter 2 (VMAT2), serotonin transporter (SERT), and norepinephrine transporter (NET) may also contribute to MPTP toxicity. It has been reported that the dopaminergic neurotoxicity of MPTP is enhanced by the injection of VMAT2 inhibitors (German et al. 2000; Reinhard et al. 1988; Staal and Sonsalla 2000) as well as in heterozygous VMAT2-knockout mice (Takahashi et al. 1997) due to the involvement of VMAT2 in the clearance of MPP<sup>+</sup> from dopaminergic neurons and the resultant accumulation of MPP<sup>+</sup> inside the cell (Liu and Edwards 1997). Additionally, the inhibition of SERT or NET by the administration of their inhibitors selectively attenuates MPTP-induced depletion of serotonin and norepinephrine, respectively (Andrews and Murphy 1993). MPTP-induced neurotoxicity is also reduced in NET-knockout mice (Rommelfanger et al. 2004). However, the functions of SERT and NET in MPTP-induced neurotoxicity still remain unknown.

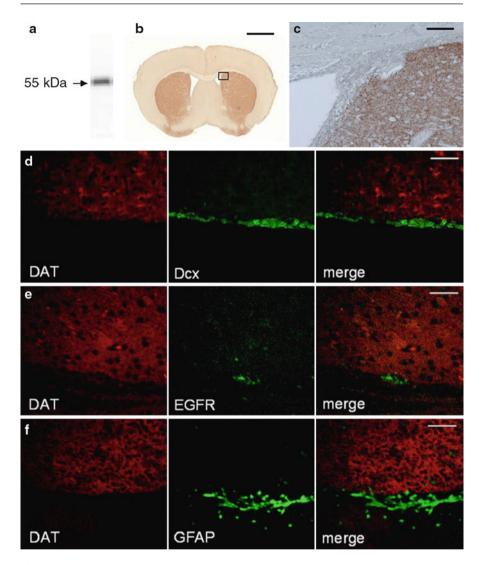
A study is performed to elucidate the role of DAT and other monoamine transporters including VMAT2, SERT, and NET on the SVZ neuroblast apoptosis induced by MPTP administration. The results of the study indicated that there are no DAT-positive neuroblasts in the SVZ (Fig. 4), whereas some neuroblasts were immunopositive for VMAT2 and SERT (Shibui et al. 2009). TUNEL-positive cells were semiquantitatively analyzed after the injection of GBR12909, a DAT inhibitor; tetrabenazine, a VMAT2 inhibitor; fluoxetine, a SERT inhibitor; or desipramine, a NET inhibitor, prior to MPTP injection. However, the injection of the transporter inhibitors had no influence on the MPTP-induced neuroblast apoptosis in the SVZ (Shibui et al. 2009). This finding suggests that the neurotoxicity of MPTP to neuroblasts in the SVZ does not require DAT or other monoamine transporters, and the MPTP-induced apoptosis may be executed through other unknown pathways.

### 4 Neurogenesis After MPTP-Induced Injury

Neurogenesis is a key event during both physiological and pathological processes, and is regulated by a variety of stimuli, such as hormones, intrinsic growth factors, neurotransmitters, exogenously applied agents, environmental enrichment, exercise, and age.

Dopamine has been shown to regulate cell cycles in the developing brain (Spencer et al. 1998). The activation of dopamine receptors influences cell proliferation in the lateral ganglionic eminence and the neuroepithelium of the frontal cortex in embryonic mice (Ohtani et al. 2003; Popolo et al. 2004). By using immunohistochemical and ultrastructural analyses, it was demonstrated that dopamine 2 like (D2L) receptors are expressed predominately on C cells in the SVZ, whereas A cells express both D1L and D2L receptors in the mouse brain (Hoglinger et al. 2004). The dopaminergic fiber is also confirmed to contact precursor cells in the SVZ of adult human and primates (Hoglinger et al. 2004; Freundlieb et al. 2006). Thus, dopamine seems to play a key role in the regulation of adult mammalian neurogenesis.

Zhao et al. (2003) first provided morphological evidences that the administration with MPTP leads to a twofold increase of bromodeoxyuridine (BrdU) incorporation in nigral dopaminergic neurons 3 weeks after the lesion in mice, which was not detectable in the granular cell layer of DG. The newly generated dopaminergic neurons are demonstrated to be derived from the cells lining the ventricular system, namely, SVZ cells. By using the nestin second intron enhancer-controlled LacZ reporter transgenic mouse model coupled with the MPTP lesion system, Shan et al.



**Fig. 4** (a) Western blot analysis demonstrates the expression of a 55 kDa DAT protein in the mouse cerebrum. (b) A representative coronal section through the striatum and SVZ were immunostained for DAT and showed the expression of DAT in the striatum, but not in the SVZ. (c) Higher magnification of the *boxed area* in (b). (d–f) The sections were also double labeled for DAT and either Dcx, EGFR, or GFAP. GFAP-positive neural stem cells, EGFR-positive neural precursor cells, or Dcx-positive neuroblasts (*green*) do not express DAT (*red*). Scale bars in (b) 1.5 mm, (c) 30  $\mu$ m, and (d–f) 15  $\mu$ m (Shibui et al. 2009)

also demonstrated that there are increased dopaminergic neurogenesis in the SN (Shan et al. 2006), supporting the findings by Zhao et al. Increased neurogenesis after the MPTP injury was also confirmed in a very recent article (Peng et al. 2008), in which MPTP lesions in the mouse brain increased the incorporation of BrdU as

well as the number of cells that co-expressed BrdU and the immature neuronal marker, doublecortin (DCX), in the DG, SVZ, and the striatum, but not in the SN, although no differentiation of newly generated cells into dopaminergic neurons was investigated. Thus, neurogenesis in the nigrostriatal system in response to MPTP damage, if it does occur, may be a later event in the self-repairing process of injured SVZ.

### 5 Strain Differences in the Susceptibility of MPTP

There are species differences in the susceptibility to MPTP injection; for example, human and monkeys are susceptible, whereas rats and hamsters are relatively resistant (Burns et al. 1983; Chiueh et al. 1984; Langston et al. 1984; Mitra et al. 1994). Mice are middle susceptible, but show marked strain differences (Sundstrom et al. 1987; Riachi and Harik 1988; Hamre et al. 1999); C57BL/6 mice are susceptible, while BALB/c mice are resistant, to MPTP neurotoxicity (Sedelis et al. 2000; Yasuda et al. 2008; Filipov et al. 2009). MPTP induces severe degeneration of dopaminergic neurons when administrated to C57BL/6 mice, but such lesions are not observed in BALB/c mice. However, the reason for such strain differences remains unclear.

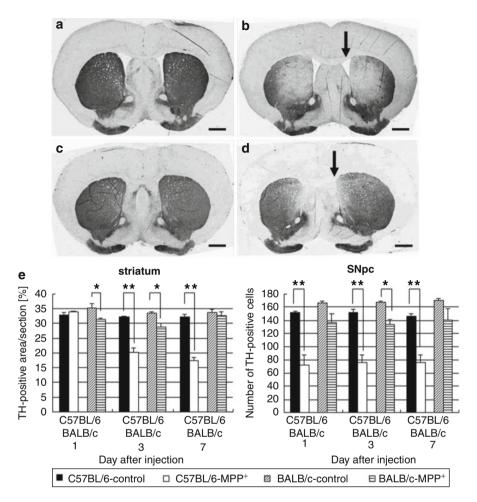
To clarify which factor(s) cause such strain differences, MPTP or MPP<sup>+</sup> was intracerebroventricularly (icv) injected into adult C57BL/6 (highly susceptible) and BALB/c (resistant) mice (Ito et al. 2013a). Both C57BL/6 and BALB/c mice injected with MPP<sup>+</sup> showed a significant decrease in tyrosine hydroxylase (TH)-immunopositive areas in the striatum and TH-positive cells in the SNpc a few days after injection, compared to saline-injected control mice, and the decrease rates in the BALB/c mice were lower than that in C57BL/6 mice. MPTP-injected C57BL/6 mice, however, showed no lesions in the striatum and SNpc (Fig. 5). These findings indicate that factors other than MAO-B can influence the strain susceptibility between C57BL/6 and BALB/c mice after the conversion from MPTP to MPP<sup>+</sup>.

Intact BALB/c mice showed higher DAT protein expression in the striatum than intact C57BL/6 mice (Ito et al. 2013b). MPTP-treated BALB/c mice showed a more significant increase of MAO-B expression than MPTP-treated C57BL/6 mice (Ito et al. 2013b). The increase of neural nitric oxide synthase (nNOS) and inducible NOS (iNOS) protein expressions in MPTP-treated BALB/c mice was more pronounced in the striatum and SNpc than in MPTP-treated C57BL/6 mice (Ito et al. 2013b).

These results indicate that MAO-B, DAT, nNOS, and iNOS expression levels do not influence the different strain susceptibility to MPTP.

### 6 Aging and MPTP Neurotoxicity

While there have been many researches on the neurotoxicity of MPTP in adult mice including our study (He et al. 2004), there have been only a few in fetal and



**Fig. 5** TH immunoreactivity in the striatum at day 7 after injection. In mice injected icv with MPP<sup>+</sup>, a reduction in TH-positive areas was observed at the injection (*arrows*) and contralateral sides of the striatum, especially adjacent to the lateral ventricles in C57BL/6 (**b**) and BALB/c (**d**) mice. However, no lesions were found in control C57BL/6 (**a**) and BALB/c (**c**) mice. Bar = 1 mm. (**e**) Strain difference in susceptibility to MPP<sup>+</sup> of C57BL/6 and BALB/c mice. Significant decreases in TH-positive areas in the striatum and TH-positive cells in the SNpc were documented in C57BL/6 mice at days 3 and 7, and at days 1, 3, and 7, respectively. In BALB/c mice, significant decreases in TH-positive areas and TH-positive cells were observed at days 1 and 3, and day 3 after injection, respectively. The decrease rates of BALB/c mice were smaller compared to those in C57BL/6 mice. Data are expressed as the mean  $\pm$  standard error (n = 3-5). \*p < 0.05 and \*\*p < 0.01 (Ito et al. 2013a)

newborn mice. Chronic MPTP exposure during the organogenesis period (days 6–15 of gestation) reduced the fetal weight, motor activity, number of striatal TH-positive cells, and amounts of DA and homovanillic acid (Wolfgang and Beat 1991). Such toxicity was enhanced when MPTP was administered postnatally

(Muthian et al. 2010). By contrast, fetal mice treated once with MPTP at gestation day 17 showed unchanged DA concentrations in the brain 24 h after injection and 24 h after birth, and in the striatum 14 and 28 days after birth compared to control mice (Melamed et al. 1990). Recently, we have demonstrated that MPTP and MPP<sup>+</sup> cause a loss of TH-positive cells and fibers and an increase of apoptotic cells in the SVZ of embryonic mice when injected into dams, and of newborn mice when injected ip (Sai et al. 2012). The results indicate that MPTP and MPP<sup>+</sup> pass through the placenta and embryonic or newborn BBB, and finally reach the brain. Further, the results of immunohistochemistry and double-labeling immunofluorescent stainings demonstrated an increase of MAO-B-positive glial cells in the brain in MPTP-treated mice, indicating the involvement of MAO-B in the acute neurotoxicity of MPTP also in both embryonic and newborn mice. The expression of DAT was not observed in the nigrostriatal area of embryonic mice and in the area and SVZ of newborn mice.

### 7 Conclusion

The administration of MPTP to susceptible animals such as C57BL/6 mice induces SVZ neuroblastic A cell apoptosis as well as SNpc and striatal dopaminergic cell injury. The conversion from MPTP to its active form, MPP<sup>+</sup>, is conducted mainly in MAO-B-positive ependymal cells and GFAP-positive cells in the SVZ. MPTP-induced neuroblast apoptosis in the SVZ is not prevented by the inhibition of monoamine transporters such as DAT, VMAT2, SERT, or NET, indicating that these transporters are not necessarily required to exert the neurotoxicity.

Recent results (Ito et al. 2012b) indicate that the nNOS or iNOS expression level is not involved in the different susceptibility to MPTP between susceptible C57BL/6 and resistant BALB/c, though there are a few reports (Chalimoniuk et al. 2006; Joniec et al. 2009) describing the significant increase of nNOS and iNOS expressions in the striatum and SNpc in C57BL/6 mice after MPTP administration. Factors influencing the strain difference in the susceptibility to MPTP toxicity are still unknown at present.

Both MPTP and MPP<sup>+</sup> can pass through the placenta and BBB and cause acute toxicity to the SN, striatum, and SVZ in the brain of embryonic and newborn mice as well as adult mice. The involvement of MAO-B in the acute neurotoxicity is confirmed regardless of the age of animals.

The replacement of lost neurons in a damaged brain area is the best way of therapeutic development for neurodegenerative diseases. Neurogenesis is, however, still a most controversial issue in studies of neurodegenerative diseases and their animal models. As mentioned, MPTP destroys neuroblastic A cells in the SVZ in the adult brain, followed by a transiently impaired neurogenesis. Though it still remains to be investigated whether and how multiple potential neural stem cells migrate into the nigrostriatal system and replace the lost dopaminergic neurons, delayed regeneration of dopaminergic neurons due to the loss of SVZ A cells may be one of possible causes of the pathological condition. With increased attention to dopaminergic neurogenesis in MPTP-induced animal models for PD, it is expected that such animals become useful as a tool for establishing the contact to PD on neurogenesis.

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### N-Methyl-(R) Salsolinol and the Enzymes Catalyzing its Synthesis and Metabolism in Parkinson's Disease

#### Makoto Naoi and Wakako Maruyama

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#### Abstract

Salsolinol (1-methyl-6,7-dihydroxy-1,2,3,4-tetrahydroisoquinoline) is an endogenous catechol isoquinoline detected in the human brain. It is produced by the

M. Naoi (🖂)

Department of Neurosciences, Gifu International Institute of Biotechnology, Kakamigahara Gifu, Japan

e-mail: mnaoi@giib.or.jp

W. Maruyama

Department of Cognitive Brain Science, National Research Center for Geriatrics and Gerontology, Obu Aichi, Japan

nonenzymatic Pictet-Spengler reaction as the racemic mixture. However, a novel enzyme, (R) salsolinol synthase, was isolated from the human brain, which catalyzes the enantio-specific synthesis of (R) salsolinol from dopamine and acetaldehyde or keto acids. R-Salsolinol is N-methylated by (R)salsolinol *N*-methyltransferase (*N*-MT) into 2(N)-methyl-(*R*)salsolinol [*N*M-(*R*)Sal], which is oxidized by an oxidase sensitive to semicarbazide. NM-(R)Salinduces cell death in the dopaminergic cells of the rat substantia nigra after the systemic administration. The behavioral and pathological changes in rats were quite similar to those in other animal models of Parkinson's disease, using 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine or 6-hydroxydopamine. In cellular experiments, NM-(R)Sal binds to type A monoamine oxidase and induces apoptosis through activation of mitochondrial death signaling. Furthermore, NM-(R)Sal levels increase in the cerebrospinal fluid from parkinsonian patients, and the activity of its synthesizing neural (R)Sal N-MT is significantly higher in the lymphocytes prepared from parkinsonian patients than in control. The enantioselective neurotoxicity of NM-(R)Sal suggests that the enzymes related to the synthesis of this toxin might be associated with the etiology for PD. This chapter reviews the role of N-methylated isoquinolines and enzymes catalyzing N-methylation of small molecules in the pathogenesis of Parkinson's disease.

#### Keywords

D	opami	ine •	Neurod	legeneration	•	Parkinson <sup>2</sup>	's	disease	• (	Salsol	lino	l
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List of Abbreviations	
CSF	The cerebrospinal fluid
DA	Dopamine
DMDHIQ <sup>+</sup>	1,2-dimethyl-6,7-dihydroxyisoquinolinium ion
IQ	Isoquinoline
IVF	Intraventricular fluid
MAO-A and MAO-B	Type A and B monoamine oxidase
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine
NM-(R)Sal	N-methyl-(R)salsolinol
N-MT	N-methyltransferase
Norsal	Norsalsolinol
PD	Parkinson's disease
ROS and RNS	Reactive oxygen and nitrogen species
Sal	Salsolinol
Sal-1-CA	Salsolinol-1-carboxylic acid
SAM	S-adenosyl-L-methionine
SN	Substantia nigra
TH	Tyrosine hydroxylase
TIQ	1,2,3,4-tetrahydroisoquinoline
β-CB	β-carboline

#### 1 Introduction

The pathological hallmark of Parkinson's disease (PD) is the selective loss of dopamine (DA) neurons in the substantia nigra (SN) and the accumulation of inclusion body called Lewy body in the brain. Several genes have been identified to be responsive for the familial form of PD. However, the molecular mechanism to develop the disease remains enigmatic, and various epidemiologic conditions have been proposed as risk factors in PD. The selective cell loss of DA neurons suggests that DA itself may be associated with the pathogenesis of PD. DA is oxidized enzymatically by monoamine oxidase [MAO, EC 1.4.3.4] and produces hydrogen peroxide and also nonenzymatically oxidized into toxic semiquinone with superoxide anion. DA oxidation induces oxidative stress, dysfunction of mitochondria and ubiquitin-proteasome system, and inflammation and activates death signal pathway in DA neurons. In addition, the discovery of 1-methyl-4-phenyl-1,2,3, 6-tetrahydropyridine (MPTP) suggested that endogenous DA-derived toxins might be involved in the neuronal loss in PD (Collins 2002).

DA-derived isoquinolines (IQs) have been proposed as endogenous neurotoxin candidates and intensively investigated in cellular and animal experiments. Salsolinol (1-methyl-6,7-dihydroxy-1,2,3,4-tetrahydroisoquinoline, Sal) is one of DA-derived 1,2,3,4-tetrahydroisoquinolines (TIQs) and was identified in urine from PD patients treated with L-DOPA (Sandler et al. 1973) and the human brain. Sal derivatives,  $\beta$ -phenylethylamine-derived TIQs, and indoleamine-derived  $\beta$ -carbolines ( $\beta$ -CBs) were found in the urine, cerebrospinal fluid (CSF), and brain (Collins 1980; Sjöquist et al. 1981). Figure 1 summarizes the chemical structure of major IQs detected in the human brain.

Racemic forms of IQs are produced by nonenzymatic condensation of monoamines with aldehydes or keto acids in the Pictet-Spengler reaction. However, the

		R <sub>1</sub>	R <sub>2</sub>	R <sub>3</sub>
HO HO R <sub>1</sub> R <sub>2</sub>	Salsolinol N-Methyl-salsolinol Norsalsolinol N-Methyl-norsalsolinol Salsolinol-1-carboxylic acid	CH <sub>3</sub> CH <sub>3</sub> H H CH <sub>3</sub>	Н Н Н СООН	H CH <sub>3</sub> H CH <sub>3</sub> H
	<ul> <li>1,2-Dehydrosalsolinol</li> <li>6- or 7-Methyl-salsolinol</li> <li>1,2-Dimethyl-6,7-dihydroxy- isoquinolinium ion</li> <li>2-Methyl-6,7-didroxy- isoquinolinium ion</li> <li>Tetrahydropapaveroline</li> </ul>	СН <sub>3</sub> Н Н Н	H H CH <sub>3</sub> H 4'-dihydroxy -benzyl)	Н Н СН <sub>3</sub> СН <sub>3</sub> Н

Fig. 1 Chemical structure of Sal derivatives identified in the human brain. \* Indicates asymmetric C<sub>1</sub> position in isoquinoline structure

predominant occurrence of the (*R*)enantiomers was confirmed in mammalian tissues (Dostert et al. 1889, 1990), suggesting the enzymatic synthesis of Sal. From the human brain, we isolated three major enzymes in Sal metabolism: (*R*)salsolinol synthase catalyzing the enantioselective synthesis *R*-Sal, a (*R*)salsolinol *N*-methyltransferase (*N*-MT) methylating *R*-Sal at 2(*N*) position into N-methyl-(*R*)salsolinol [*NM*-(*R*)Sal], and *NM*-(*R*)Sal oxidase producing 1,2-dimethyl-6, 7-dihydroxy-1,2,3,4-tetrahydroisoquinoline ion (DMDHIQ<sup>±</sup>).

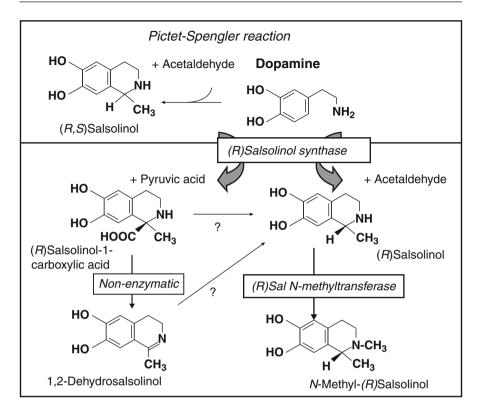
This chapter summarizes the results on the enzymatic biosynthesis and metabolism of (*R*)Sal and *N*M-(*R*)Sal, the occurrence of Sal derivatives in the human brain, the preparation of in vivo model of PD in rat, and the molecular mechanism behind the programmed cell death induced by *N*M-(*R*)Sal. Finally the analyses of clinical samples from PD patients are presented to show the possible association of *N*M-(*R*)Sal with the pathogenesis of PD. The role of *N*M-(*R*)Sal and a neutral (*R*)Sal *N*-MT and other *N*-MTs in the selective degeneration of DA neurons in PD is discussed.

#### 2 Biosynthesis of (R)salsolinol Derivatives in Human Brain

#### 2.1 (R)Salsolinol Synthase Catalyzing the Enantio-Specific Synthesis of (R)salsolinol

In foods and beverages, Sal occurs as racemic mixture produced by the nonenzymatic condensation in the Pictet-Spengler reaction. However, Sal has an asymmetric center at 1 position, and (R)-enantiomer was dominant in urine from healthy volunteers, suggesting the stereoselective biosynthesis (Dostert et al. 1989). Dostert et al. (1990) proposed the biosynthesis pathway of (R)Sal in human, as shown in Fig. 2. The condensation of DA with pyruvic acid yields salsolinol-1-carboxylic acid (Sal-1-CA), which was detected in the human brain caudate nucleus, the CSF, and urine, but the enantiomeric structure was not determined (Ung-Chhun et al. 1985). The nonenzymatic decarboxylation of Sal-1-CA produces 1,2-dehydrosalsolinol, which was also identified in human urine (Dostert et al. 1990). The presence of an enzyme catalyzing the oxidative decarboxylation of Sal-1-CA into 1,2-dehydrosalinol was reported in rat kidney (Collins and Cheng 1988), but the enzyme has neither been characterized nor detected in the human brain. The enantioselective reduction of 1,2-dehydrosalsolinol or the direct decarboxylation of Sal-1-CA into (R)Sal has never been confirmed in animals or plants.

The enzyme participating in the enantio-specific biosynthesis of (R)Sal was isolated from the human brain. (R)Sal synthase was purified from the cytosolic fraction prepared from human gray matter as a monomer with molecular mass of 34.3 kDa. Table 1 summarizes the basic characteristics of this enzyme. (R)Sal synthase catalyzes the enantioselective synthesis of (R)Sal and (R)Sal-1-CA from DA with acetaldehyde or pyruvic acid (Naoi et al. 1996b). This enzyme synthesizes also norsalsolinol (6,7-dihydroxy-1,2,3,4-tetrahydroisoquinoline, Norsal) from DA and formaldehyde, but adrenalin and noradrenalin are not the substrate.



**Fig. 2** Nonenzymatic and enzymatic synthesis of (*R*)-enantiomer of Sal derivatives in the human brain. Dopamine is condensed with acetaldehyde into (*R*/S)Sal by the Pictet-Spengler reaction. (*R*) Salsolinol synthase condenses DA with acetaldehyde or pyruvic acid to yield (*R*)Sal and (*R*)Sal-1-carboxylic acid, which is nonenzymatically converted into1,2-dehydrosalsolinol. The enantioselective synthesis of (*R*)Sal from (*R*)Sal-1-carboxylic acid or 1,2-dehydrosalsolinol has not been confirmed. SAM-dependent *N*-methyltransferase catalyzes the *N*-methylation of (*R*)Sal into *N*M-(*R*)Sal. *N*M-(*R*)Sal is further oxidized into DMDHIQ<sup>+</sup> enzymatically by *N*M-(*R*)Sal oxidase or nonenzymatically

The enzymatic activity to produce Norsal is the highest, followed by that to (R)Sal-1-CA and (R)Sal. However, this synthase does not produce NM-(R)Sal directly from N-methyldopamine (epinine) with acetaldehyde, suggesting that NM-(R)Sal is synthesized by two-step reactions with (R)Sal synthase and an N-MT.

#### 2.2 (R)Salsolinol N-Methyltransferase

The conversion of N-methylation of (R)Sal into 132 NM-(R)Sal was con rmed in our studies using in vivo microdialysis in rat brain, and the highest activity was in the SN (Maruyama et al. 1992). In the sample from the human brain, (*R*)Sal *N*-MT activity was detected with two peaks at pH 6–7 and 8–9, and the activity of sample prepared from the striatum was higher at pH 7.0 than at the more alkaline pH

	(R)salsolinol synthase	Neutral ( <i>R</i> )salsolinol <i>N</i> -Methyltransferase
Substrate specificity	Dopamine, acetaldehyde, pyruvic acid	(R)salsolinol, norsalsolinol
	Not <i>N</i> -methyl-DA, adrenaline, noradrenaline	
Methyl donor		SAM
Optimal pH	7.4	7.0
Molecular weight	34.3 kDa <sup>a</sup>	35.6 kDA <sup>b</sup>
Intracellular localization	Cytoplasm	Cytoplasm
Michaelis constant (Km)	For dopamine; $5.11 \pm 0.62 \text{ mM}$	For ( <i>R</i> )Sal; $126 \pm 73 \mu\text{M}$
	For acetaldehyde; $0.59 \pm 0.12 \text{ mM}$	For SAM; 57.6 $\pm$ 2.3 $\mu M$
Enzymatic activity	( <i>R</i> )Sal; $342.4 \pm 40.5$	<i>N</i> M-( <i>R</i> )Sal; $28.4 \pm 13.6$
(pmol/min/mg protein)	( <i>R</i> )Sal-1-carboxylic acid; 1 149 $\pm$ 95	NM-Norsal; $9.1 \pm 3.0$
	Norsal; 3 715 $\pm$ 207	

Table 1 Characteristics of enzymes related to the metabolism of *N*-methyl-(*R*)salsolinol

(R)Sal synthase and (R)Sal *N*-methyltransferase were isolated from the cytoplasm fraction of human brain gray matter and the striatum, respectively

 $^{a}(R)$ Sal synthase was purified by stepwise column chromatograph with twice large DEAE, once small DEAE, and finally gel-filtration column, to a single protein with 1 700-fold purification from the cytoplasm fraction (Naoi et al. 1996a, b)

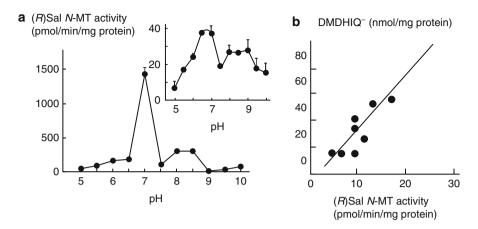
<sup>b</sup>(*R*)Sal *N*-methyltransferase was purified by stepwise column chromatograph with a DEAE, and finally gel-filtration column, to a single protein with 130-fold purification from the cytoplasm fraction

(Fig. 3a, inserted). A neutral (*R*)Sal *N*-MT with the optimum pH at 7.0 was isolated from the cytoplasm fraction of the human brain as a single protein with the molecular mass of 35.6 kDa (Fig. 3a) (Naoi et al. 1997). *S*-Adenosyl-L-methionine (SAM) was a methyl donor. This neutral *N*-MT had the strict affinity to (*R*)Sal, and (*S*)Sal was not the substrate. Previously, a *N*-MT for 1,2,3,4-TIQ was isolated from the human brain, and the reaction was optimal at pH 8.25 (Naoi et al. 1989), which might correspond to alkaline (*R*)Sal *N*-MT. There was no correlation between the activities of the neutral and alkaline *N*-MT, and the age and sex did not affect the activities in any brain regions (Maruyama et al. 1997c).

Amine *N*-MT A from bovine liver was reported to *N*-methylate stereospecifically (*R*)-norsalsoline (7-hydroxy-6-methoxy-1-methyl-1,2,3,4-TIQ) and (*R*)salsolidine (6,7-dimethoxy-1-methyl-1,2,3,4-TIQ) (Bahnmaier et al. 1999). Recombinant human amine *N*-MT [tryptamine *N*-MT, EC 2.1.1.49] concerts benzylisoquinoline, tetrahydropapaveroline (THP), 6-O-methyl-THP and norreticuline into the *N*-methylated derivatives with a strict preference for (*R*)-configuration (Grobe et al. 2011). At present, it is not confirmed whether they can *N*-methylate Sals.

## 2.3 *N*-Methyl-(*R*)salsolinol Oxidase and Other Enzymes Related to the Metabolism

An enzyme catalyzing the oxidation of NM-(R)Sal into DMDHIQ<sup>±</sup> was identified in the human brain (Naoi et al. 1995). This oxidase is sensitive to semicarbazide,



**Fig. 3** The pH-dependent activity of (*R*)Sal *N*-MT and relation to Sal derivatives in the human brain. (a) The effect of pH on neutral (*R*)Sal *N*-MT activity purified from the human striatum. Inserted figure represents the effect of pH on the activity of (*R*)Sal *N*-MT in the cytoplasm fraction for the human striatum. (b) The relationship of DHDMIQ<sup>+</sup> levels in the SN with neutral (*R*)Sal *N*-MT activity in the caudate from the control human brains. Significant correlation was confirmed between the DMDHIQ<sup>+</sup> level in the substantia nigra and the activities of neutral *N*-methyltransferase in the caudate (r = 0.905, p < 0.001) and putamen (r = 0.616, p < 0.05)

but not inhibited by clorgyline and deprenyl, inhibitors of type A and B monoamine oxidase (MAO-A, MAO-B). It is localized in cytoplasm, but not in mitochondria as in the case with MAO. It catalyzes also the oxidation of NM-(S)Sal and 2(N)-methyl Norsal (NM-Norsal) with much less activity than to NM-(R)Sal. (R)- and (S)-Sal and Norsal were not the substrate of this enzyme. NM-(R)Sal is also nonenzymatically oxidized (Ung-Chhun et al. 1985; Maruyama et al. 1995).

#### 3 The Occurrence of Salsolinol Derivatives in the Human Brain

#### 3.1 Occurrence of (R)- and (S)-Enantiomers

The reported contents of Sal derivatives in the human brain basal nuclei were varied from 0.2 to 10 ng/g wet tissue weight, indicating the difficulty in the quantitative analyses of human materials. The occurrence of (R)- and (S)-Sal derivatives in the human brain was reported, but the results are contracting, as summarized in Table 2. We quantitatively determined (R)- and (S)-Sal derivatives in the human brain from control without the history of any neurological disorder (Maruyama et al. 1997a). (R)Sal was detected ubiquitously in the frontal lobe, caudate, putamen, and SN, but the (S)Sal was under detection limit. NM-(R)Sal content was significantly high in the SN and DMDHIQ<sup>+</sup> was detected only in the SN. Others reported similar results. Sal derivative contents were high

· · ·			
Reference	Tissue, brain regions	Isoquinolines	Enantiomer
Moser and Kämpf (1992)	CSF	Sal, NM-Norsal	
Moser et al. (1995)	CSF	NM-Norsal <sup>a</sup>	
Maruyama et al. (1996a)	CSF	Sal, NM-Sal <sup>b</sup>	only (R)-
Maruyama et al. (1999)	CSF	Sal, NM-Sal <sup>a</sup>	only (R)-
Maruyama et al. (1996b)	IVF	<i>N</i> M-Sal <sup>a</sup> , (Sal), DMDHIQ <sup>+</sup>	only (R)-
Maruyama et al. (1997c)	Frontal lobe, caudatus, putamen, substantia nigra	Sal, NM-Sal <sup>a</sup>	only $(R)$ -
Müller et al. (1999)	CSF	(R), (S)-Sal <sup>c</sup>	(R) = (S)
Musshoff et al. (1999)	Hypothalamus, accumbens, caudatus, putamen	Sal, Norsal	Not determined
Musshoff et al. (2000)	N. accumbens, caudatus, putamen, SN, hypothalamus	Sal, Norsal	(R) > S (R/S = 2)
DeCuypere et al. (2008a)	Putamen, substantia nigra,	Sal <sup>a</sup> , <i>N</i> M-Sal <sup>a</sup> , Norsal	(R) = (S)
	Cerebellar cortex, frontal lobe		R > S
DeCuypere et al. (2008b)	SN, striatum	NM-Norsal <sup>a</sup>	

**Table 2** Occurrence of (R)- and (S)-enantiomers of Sal derivatives in the human brain, CSF, and IVF (Reported later than 1990)

<sup>a</sup>Decreased, <sup>b</sup>increased, or <sup>c</sup>not changed in PD and aging

in the striatum followed by the SN and hypothalamus (Musshoff et al. 1999; DeCuypere et al. 2008a). However, Musshoff et al. (1999) reported the presence of minor amounts of (S)Sal in addition to (R)Sal with ration of about 1–2, and DeCuypere et al. (2008a) found the same amounts of (R)- and (S)-enantiomers of Sal and NM-Sal in the caudate and putamen, SN, and frontal lobe. At present, the reason of the discrepancy has not been well clarified. However, it is clear that (R)-enantiomers of Sal and NM-Sal are synthesized in the brain, and the enzymatic synthesis of (R)Sal derivatives is predominant in the brain regions with low DA contents, suggesting that nonenzymatic synthesis of racemic Sal might occur in the regions containing high levels of free DA. In addition, Sal derivatives may be transported into the brain through the blood–brain barrier (BBB) (Thümen et al. 2002), and racemic Sal derivatives derived from food and beverages might contribute to the presence of (S)Sal derivatives in the brain.

In the human brain, Norsal was also detected in the striatum, followed by the SN and hypothalamus (Musshoff et al. 1999). *NM*-Norsal was identified also in the same regions and immunochemically co-localized with a neuronal marker, neuron-specific enolase, and a catecholamine marker, tyrosine hydroxylase (TH) (DeCuypere et al. 2008b).

In neurons, Sal derivatives are localized in distinct intracellular compartments depending on their chemical structure. Sal derivatives are accumulated in the mitochondrial and membrane-nuclear fraction, 90–60 % of the total amounts, after 3-day incubation of (*R*)Sal and Norsal with PC12 cells (Maruyama et al. 1993). These results might be relevant with the recent result that *N*M-(*R*)Sal binds to MAO-A in the mitochondria (Yi et al. 2006). DMDHIQ<sup>+</sup> binds to melanin, which may cause its high contents in the SN (Naoi et al. 1994a). <sup>3</sup>H-Sal was reported to bind to a selective binding site in the striatum, cortex, median eminence, and hypothalamus of the rat brain, and the binding was displaced with DA, L-DOPA, carbidopa, and apomorphine, but not by DA receptor ligands, inhibitors of DA transport (Homicsko et al. 2002). The presence of this Sal-binding protein has not been confirmed in the human brain.

#### 3.2 Factors Affecting the Salsolinol Levels

In human brain samples, (*R*)Sal levels decreased in the caudate, putamen, and SN according to the age, and a negative correlation was confirmed between NM-(*R*)Sal levels in the striatum and the age (Maruyama et al. 1997c). Alcohol intake markedly increased racemic Sal in blood and maybe also in the brain, but this issue has not been settled. The presence of alcohol did not affect the levels of (*R*)-enantiomer of Sal and *N*M-Sal in any brain regions (Naoi et al. 2004).

These results suggest that the concentration of the (R)-enantiomers may depend on the activity of the synthesizing enzymes. The relationship between the activities of Sal-related enzymes and Sal derivative levels was examined in human brain regions (Naoi et al. 1997). A good correlation was confirmed between the activity of neutral (R)Sal N-MT, but not that of alkaline N-MT, in the striatum (putamen and caudate) and the concentration of DMDHIQ<sup>+</sup> in the SN (Fig. 3b). NM-(R)Sal is synthesized in the striatum, oxidized into DMDHIQ<sup>+</sup>, and transported and accumulated in the SN, as in the case with the rat PD model, where DMDHIQ<sup>+</sup> was detected in the SN after injection of NM-(R)Sal in the striatum (Naoi et al. 1996b). There was no other relationship that was confirmed between other enzyme activities and Sal derivative levels in frontal cortex and thalamus. In rat brain, the levels of racemic NM-Sal reached to the maximum within 2–6 h after the intraperitoneal injection, but NM-Sal was quickly eliminated from the brain (Lorenc-Koci et al. 2008). This rapid clearance of Sal may account for the discrepancy of the Sal contents with the activity of synthesizing enzymes in the human brain regions.

Sal derivatives are O-methylated by catechol-O-methyltransferase (COMT) (Hötzl and Thomas 1997) and oxidized by FAD-containing monooxygenase (Wu et al. 1992) and debrisoquine 4-monooxygenase (Iwahashi et al. 1993). However, it has not been studied whether the same enzymes are associated with the catabolism of Sal derivatives in the human brain.

#### 4 Neurotoxicity of *N*-Methyl-(*R*)salsolinol: In Vivo and In Vitro Studies

#### 4.1 An Animal Model of Parkinson's Disease

A PD model was established with NM-(R)Sal. After 1-week continuous injection in the rat striatum, NM-(R)Sal caused behavioral changes, such as akinesia, twitching, postural disturbance, and spontaneous rotation (Naoi et al. 1996b). The number of DA neurons stained with TH antibody and the contents of DA, noradrenaline, and their metabolites reduced markedly in the SN of the injected site. On the other hand, NM-(S) Sal, (R)Sal and (S)Sal, Norsal, and NM-Norsal did not induce significant biochemical-histopathological changes in rats. DMDHIQ<sup>+</sup> caused severe necrotic reaction at the injected site, but the behavioral and neurochemical changes were only transit.

The preparation of animal models of PD with Sal and other endogenous IQs has been also reported. Chronic administration of racemic Sal and TIQ caused a decrease in DA metabolism in the SN and striatum of rats, suggesting the loss of DA neurons (Antkiewicz-Michaluk et al. 2000). Systematic administration of racemic *N*M-Sal into rat perturbed CA metabolism but failed to cause the cell death in the SN (Lorenc-Koci et al. 2008). TIQ and *N*-methyl-6,7-dimethoxyisoquinolinium ion were infused in rat supra-nigral region and produced 42 % and 20 % SN cell loss, respectively, and reduced DA contents in the striatum, but behavioral change was not observed (McNaught et al. 1996). Intraperitoneal injection of TIQ reduced TH-immunoreactive neurons and DA levels in the striatum (Lorenc-Koci et al. 2000). Administration of indoleamine-derived 2(N)-methyl- $\beta$ -CB into the rat SN induced the reduction of DA metabolites and the massive necrotic features (Neafsey et al. 1989). The presence of a methyl group at 9(N) position of  $\beta$ -CB increased the in vivo neurotoxicity markedly (Matsubara et al. 1998).

## 4.2 Induction of Apoptosis by *N*-Methyl-(*R*)salsolinol and Its Molecular Mechanism

In PD, apoptosis of dopamine neurons was confirmed in the SN by detection of activated caspase 3, increased Bax, and nuclear translocation of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (Hartmann et al. 2000; Tatton 2000). In human dopaminergic neuroblastoma SH-SY5Y cells, *NM*-(*R*)Sal induces apoptosis (Akao et al. 1999, 2002; Maruyama et al. 1997b, 2001a, b). Among Sal derivatives, *NM*-(*R*) Sal is the most potent to induce DNA damage, whereas *NM*-(*S*)Sal, (*R*)- and (*S*)-Sal, and 1-carboxy-Sal are much less cytotoxic. Apoptosis is initiated with increased membrane permeability in mitochondria, as shown by the collapse in membrane potential,  $\Delta\Psi$ m, and swelling of mitochondria. Then, cytochrome c is released in the cytoplasm, caspase 3 is activated, GAPDH is translocated in nuclei, and finally nucleosomal DNA is fragmented and condensed. Only (*R*)-enantiomer of *NM*-Sal induced  $\Delta\Psi$ m collapse in SH-SY5Y cells (Maruyama et al. 2001a, b) and also in isolated mitochondria (Akao et al. 2002), suggesting the presence of a binding protein of *NM*-(*R*)Sal. *NM*-(*R*)Sal binding to MAO-A located in mitochondria

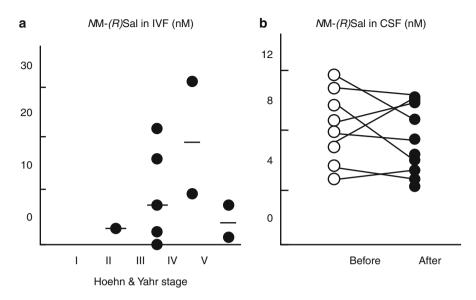
outer membrane initiates apoptosis signaling (Yi et al. 2006). *NM*-(*R*)Sal binds to MAO-A at the substrate-binding site, which is suppressed by knockdown of MAO-A expression by short-interfering RNA (siRNA) and also by serotonin (5-hydroxytryp-tamine, 5-HT) and clorgyline, a MAO-A-specific substrate and inhibitor.

Sal derivatives cause also oxidative stress and mitochondrial dysfunction, proposed causal factors for dopaminergic neural loss in PD. *NM*-(*R*)Sal increases reactive oxygen and nitrogen species (ROS, RNS) in the dialysate of in vivo microdialysis at the rat striatum, but (*R*)Sal reduces the ROS levels at the distinct concentrations (Maruyama et al. 1995). (*R*)Sal, *NM*-(*R*)Sal, and DHDMIQ<sup>+</sup> inhibit the activities of complexes I (NADH-Q reductase) and complex II (succinate-Q reductase) in mitochondrial electron transfer pathway, and DHDMIQ<sup>+</sup> is the most potent inhibitor (Morikawa et al. 1998). *N*-Methylation of Sal and TIQ increases the inhibition of complex I and the cytotoxicity, as shown with *NM*-isoquinolinium ion (*NM*-IQ<sup>+</sup>) and *NM*-TIQ (Suzuki et al. 1992), 1,2,3,4-tetrahydro-2-methyl-4,6,7isoquinolinetriol (Willets et al. 1996), and *NM*-Norsal (Storch et al. 2002). Increased ROS production plays a major role in the cytotoxicity of racemic Sal, which particularly inhibits complex II (Storch et al. 2000). TIQ and IQ induced apoptosis in PC12 and SK-N-MC cells through the inhibition of complex I and increased ROS production (Seaton et al. 1997).

#### 5 *N*-Methyl-(*R*)salsolinol and *N*-Methyltransferases in Parkinson's Disease

#### 5.1 Salsolinol Derivatives in the Parkinsonian Brain

The changes in the amounts of Sal derivatives have been scarcely reported in the brain and CSF from PD patients. In the caudate nuclei of the brains from PD patients, the lower levels of (R)Sal, (S)Sal, NM-(R)Sal, and NM-(S)Sal were reported in comparison with the normal brain (DeCuypere et al. 2008a). On the other hand, in the CSF, NM-(R)Sal was increased in PD compared to control and patients with multiple system atrophy (MSA) (Maruyama et al. 1996a). The NM-(R)Sal contents in PD (n = 16) were 8.32  $\pm$  2.89 nM (4.54–15.7), whereas in control (n = 29) and MSA (n = 5)  $4.53 \pm 2.08$  (0.62–11.9) and  $3.59 \pm 1.52$ (1.33-5.25). The relative concentration of NM-(R)Sal versus homovanillic acid (HVA), a DA metabolite, is significantly higher in PD than in control and MSA, suggesting the increased synthesis of NM-(R)Sal in PD. In the PD CSF, Sal was detected (Moser and Kämpf 1992) and Sal was increased in PD patients with dementia (Antkiewics-Michaluk et al. 1997). In the intraventricular fluid (IVF) from parkinsonian patients treated with L-DOPA, NM-(R)Sal and a trace amount of (R)Sal were detected in addition to DMDHIQ<sup>+</sup>, but the (S)-enantiomers were not (Maruyama et al. 1996b). The amounts of NM-(R)Sal in the CSF and IVF had a tendency to increase at the early Hoehn and Yahr stage and decrease according to the advancing Hoehn and Yahr stage by disease duration (Maruyama et al. 1999) (Fig. 4a). In the CSF from the PD patients (n = 9), NM-R-Sal levels



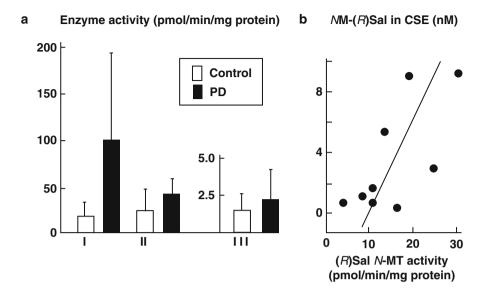
**Fig. 4** The effects of Hoehn and Yahr stage and disease duration on NM-(R)Sal levels in the IVF and CSF. (a) The effects of the Hoehn and Yahr stage on the NM-(R)Sal levels in the IVF from parkinsonian patients. (b) Decline of NM-(R)Sal levels in the CSF of the same PD patients after 2 years of disease duration

decreased after 2 years of L-DOPA medication (Fig. 4b). The levels of Sal and NM-Norsal in the CSF were negatively correlated with the duration of PD (Moser et al. 1995; Müller et al. 1999). In plasma from de novo parkinsonian patients, (R)- and (S)-Sal levels were not increased, but (R)Sal levels were inversely related to the duration and intensity of PD, in contrast to (S)Sal and DA (Müller et al. 1998), whereas L-DOPA therapy increases Sal, Norsal, and NM-Norsal in urine (Scholz et al. 2004).

These results indicate that Sal derivative increase at the early phase of PD and reduce according to the disease progression. The loss of DA neurons caused by the disease progression and also aging reflects the Sal levels in the postmortem brain samples and in the CSF and IVF. Significant reduction of NM-(R)Sal in the SN for PD brains compared to control (DeCuypere et al. 2008a) may indicate the accelerated cell loss of DN neurons by this neurotoxin.

#### 5.2 Neutral (R)salsolinol and Other N-Methyltransferases

The enzyme activities for Sal synthesis and metabolism were investigated in lymphocytes isolated from PD patients (n = 24) and control subjects without neurological disease (n = 56) (Naoi et al. 1998). The enzymatic activities of neural (*R*)Sal *N*-MT were found to increase significantly in PD lymphocytes, whereas the activities of alkaline (*R*)Sal *N*-MT and *N*M-*R*-Sal oxidase were not changed (Fig. 5a). The relationship between (*R*)Sal *N*-MT activity in the lymphocytes and

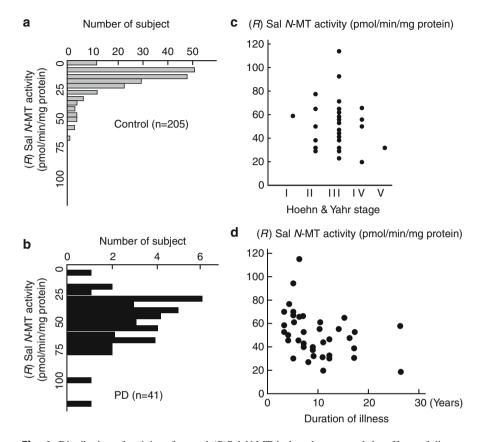


**Fig. 5** The activities of (*R*)Sal *N*-MT and related enzymes in lymphocytes from PD patients and its relation to NM-(*R*)Sal levels in the CSF. (**a**) The enzymatic activity of neutral (I) and alkaline (*R*)Sal *N*-MT and *NM*-(*R*)Sal oxidase (III) in lymphocytes prepared from control (*white column*) and PD patients (*black column*). The column and bar represent the mean and SD. (**b**) The relationship of neutral (*R*)Sal *N*-MT activity in lymphocytes with *NM*-(*R*)Sal levels in the CSF from the same PD patients

*N*M-(*R*)Sal levels in the CSF was investigated (Fig. 5b). A weak correlation between them was observed (y = -3.65 + 0.62, R = 0.71, p < 0.05), but the limited number of the samples (n = 9) could not yield a conclusive result.

The increase in (*R*)Sal *N*-MT in PD lymphocytes was reexamined in other groups of control and PD patients (Fig. 6a, b). The activities of (*R*)Sal *N*-MT in PD lymphocytes (n = 41) were mainly distributed as being higher than 35 pmol/min/ mg protein, whereas in control lymphocytes (n = 205) the activities were mostly lower than 30 pmol/min/mg protein. The activities did not show significant relationship to Hoehn-Yahr stage and the duration of illness, in contrast to the *N*M-(*R*)Sal contents in the CSF and IVF (Fig. 6c, d). The activities in lymphocytes were not affected by the gender, the age of PD onset, and the age of the patients when the samples were taken, and the dose of L-DOPA administered. These results suggest that (*R*)Sal *N*-MT activity may be genetically determined and the increased activity observed in PD may be neither secondary to the disease process nor to the treatment. (*R*)Sal *N*-MT was isolated and purified from the control brain, but we have no chance to investigate whether this enzyme changes its enzymatic properties in the PD brain.

*N*-Methylation of Sal, TIQs, and  $\beta$ -CB increases the cytotoxicity, as discussed above (Naoi et al. 1994b; Maruyama et al. 1997b; Matsubara et al. 2002; Williams and Ramasden 2005). Mammalian SAM-dependent *N*-MT [EC. 2.1.1] is a large family of enzymes catalyzing *N*-methylation of small molecules, and these *N*-MTs have similar molecular weights in the range of 25–35 kDa and share common enzymatic



**Fig. 6** Distribution of activity of neutral (*R*)Sal *N*-MT in lymphocytes and the effects of disease progress. (a) The activity in control subjects (n = 205). (b) In lymphocytes from PD patients (n = 41), the activity was distributed to higher than 35 pmol/min/mg protein. (c) The effects of the Hoehn and Yahr stage on the activity of neutral (*R*)Sal *N*-MT in lymphocyte from PD patients. (d) The effects of disease duration on the activities of neutral (*R*)Sal *N*-MT in lymphocytes from PD patients

characteristics. Nicotinamide *N*-MT [EC 2.1.1.1] is isolated from human liver, has a molecular weight of 29kDa as a monomer, and *N*-methylates nicotinamide and other pyridine (Aksoy et al. 1994). This *N*-MT converts 4-phenyl-1,2,3,6-tetrahydropyridine and 4-phenylpyridine into MPTP and MPP<sup>+</sup>, suggesting the biosynthesis of MPTP derivatives from endogenous or environmental precursors (Ansher et al. 1986). The *N*-methylation of nicotinamide produces *N*-methylnicotinamide, which has MPP<sup>+</sup>like cytotoxicity (Fukushima et al. 2002). The protein levels of nicotinamide *N*-MT were higher in the CSF from PD patients (65 years old or younger) than control and decreased with aging (Aoyama et al. 2001). In the cerebella and caudate from PD patients, high expression of this *N*-MT protein and activity was confirmed, and the disease duration was negatively correlated with the expression (Parsons et al. 2002). The increased nicotinamide *N*-MT in PD may be relevant with the increased levels of *N*-methylnicotinamide in parkinsonian brain (Williams et al. 1993) and high urinary excretions of nicotinamide metabolites in younger PFD patients, which declined with aging (Aoyama et al. 2000).

In PD brain, the activity of  $\beta$ -CB 9(*N*)-MT, which methylates nitrogen at 9 position in  $\beta$ -CB, but not at 2, increased in the frontal cortex in PD (Gearhart et al. 2000). Phenylethanolamine (PEA) *N*-MT (noradrenalin *N*-MT), which catalyzes the methylation of noradrenalin into adrenalin [EC 2.1.1.28], was reported to have  $\beta$ -CB 2(*N*)-MT, which methylates nitrogen at 2 position of  $\beta$ -CB (Gearhart et al. 2002). The thr15Ile polymorphism of a histamine *N*-MT [EC 2.1.1.8] was reported to be associated with PD (Palada et al. 2011) and the increase in histamine, which is related to DA neuron loss in PD (Anichtchik et al. 2000). However, recent papers did not confirm the association of this functional polymorphism with PD and essential tremor (Keeling et al. 2010) and AD (Marasovic-Susnjara et al. 2011).

#### 6 Conclusion

These presented results from the basic research using cellular and animal models to the analyses of clinical samples from parkinsonian patients prove that NM-(R)Sal may be involved in the pathogenesis of PD as an endogenous dopaminergic toxin. However, these results cannot be free from criticism. The quite high concentrations of these IQs are used for very short time of experiment, which might not reasonably reflect the slowly processing neuronal loss in PD. The postmortem studies of IQ contents in the parkinsonian brain cannot draw the common consensus because of unavoidable variations in the quantitative data. These results seem to suggest that the neurotoxin hypothesis might pass away as the pathogenesis of PD, if we cannot establish new approach.

However, the results on (*R*)Sal *N*-MT, nicotinamide *N*-MT, and other *N*-MT suggest that we should investigate the enzymes in the metabolism of Sal derivatives in relation to the pathogenesis of PD. *N*-MT might be a good candidate for this study, since this enzyme can convert small molecules of endogenous and environmental protoxins into cytotoxic *N*-methylated toxins. The activity is regulated genetically and also by environmental factors, protoxin levels in diets, and cellular signals, such as interferon- $\gamma$  (Yamada et al. 2010). These results suggest that the study on *N*-MT brings us further insights on the genetic and environmental factors as the pathogenesis of PD. If we can find the methods to suppress the synthesis of toxins with inhibitors, we can develop novel neuroprotective therapy to prevent the onset and progress of degeneration of DA neurons.

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Declaration of Approval from the Ethical Committee

The Ethical Committee of Iwate Medical University, Aichi Medical University, National Institute for Longevity Science, and Neurological Clinic (Dr. H. Narabayashi) approved the protocol for the examination of the CSF samples and lymphocytes and of the IVF. The patients were fully informed about the risk and the benefits of the examination.

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# Regulation of DA Homeostasis and Role of VMAT2 in DA-Induced Neurodegeneration

Eugene V. Mosharov

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#### Abstract

Metabolic turnover of dopamine (DA) and other monoamines is tightly regulated by their synthesis, degradation, and compartmentalization. DA transmission plays important roles in learning and behavior, while abnormal DA tone is implicated in multiple neurological disorders, including Parkinson's disease (PD), schizophrenia, and psychoses. Free cytosolic DA can produce oxidative stress and protein damage and is suspected to contribute to the development of PD. Vesicular monoamine transporter 2 (VMAT2) sequesters DA into synaptic vesicles, thereby occupying a unique position where it facilitates DA synaptic transmission while preventing the deleterious effects of DA presence in the cytosol. This review summarizes recent findings on the regulation of DA cellular homeostasis and on the relationship between VMAT2 activity and DA-mediated neurotoxicity.

E.V. Mosharov

Department of Neurology, Columbia University, New York, NY, USA e-mail: em706@columbia.edu

#### Keywords

Dopamine • Neurotoxicity • VMAT

List of Abbreviations					
AADC	Aromatic L-amino acid decarboxylase				
DA	Dopamine				
DA <sub>cyt</sub>	Cytosolic DA concentration				
DAT	DA uptake transporter				
DOPAC	3,4-dihydroxyphenylacetate				
DOPAL	3,4-dihydroxyphenylacetaldehyde				
LC	Locus coeruleus				
L-DOPA	L-3,4-dihydroxyphenylalanine, levodopa				
MAO	Monoamine oxidase				
METH	Methamphetamine				
$MPP^+$	1-methyl-4-phenylpyridinium				
MSN	Medium spiny neurons				
PD	Parkinson's disease				
SNpc	Substantia nigra pars compacta				
ROS	Reactive oxygen species				
TH	Tyrosine hydroxylase				
VGLUT2	Vesicular glutamate transporter 2				
VTA	Ventral tegmental area				
VMAT	Vesicular monoamine transporter				

#### 1 Introduction

Dopaminergic transmission in the brain plays important role in movement control, decision making, motivation, reward, learning, and other essential behavioral and cognitive functions. The three most thoroughly characterized dopaminergic projections are the nigrostriatal, mesolimbic, and mesocortical systems that each originate in the ventral midbrain and project to the striatum, limbic system, and cortex, respectively. These systems are targeted by drugs of abuse and psychostimulants, and abnormal DA neurotransmission in these pathways is linked to a number of diseases. Altered sensitivity to DA in mesolimbic and mesocortical pathways has been implicated in schizophrenia, narcolepsy, and effective disorders (van Rossum 1966; Gamo and Arnsten 2011), whereas degeneration of nigrostriatal neurons in PD represents the best-studied link between DA neurotransmission and neuropathology (Fahn 2008; Obeso et al. 2008).

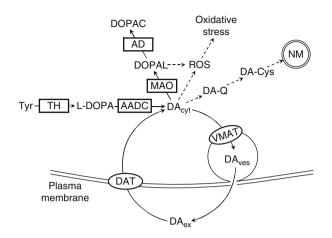
PD is characterized by the preferential loss of DA neurons in the substantia nigra pars compacta (SNpc), although other neuronal populations, including locus coeruleus (LC), raphe nuclei, the dorsal motor nucleus of the vagus nerve, and the peripheral autonomic neurons, are also affected. Neurodegeneration in SNpc with the concomitant depletion of striatal DA is responsible for the cardinal motor symptoms of PD: tremor, rigidity, bradykinesia, and postural instability (Fahn 2008; Obeso et al. 2008). Other features of PD include inflammation and the appearance of alpha-synuclein-positive "inclusions" known as Lewy bodies and Lewy neurites. Age is the greatest risk factor of PD; the likelihood that an individual will develop the disease is ~0.5 % at the age of 65 and increases by ~0.2 % each subsequent year (de Rijk et al. 1997; Baldereschi et al. 2000; Fahn and Sulzer 2004). The majority of PD cases are idiopathic, although at least 10 % of patients are now thought to have "familial" forms of the disease (reviewed in Martin et al. 2010); research to unveil new genetic causes of PD still continues.

SNpc and LC contain large quantities of neuromelanin, a dark pigment that gives names to these brain areas. Neuromelanin appears in SNpc neurons within 3 years of birth and increases linearly with age (Zecca et al. 2002). The pigment is derived from the oxidized metabolites of DA that form stable complex with iron, copper, and other metals (Sulzer et al. 2000; Zucca et al. 2006) and can be produced by DA neurons in vitro within several days of treatment with DA precursor L-3,4-dihydroxyphenylalanine (L-DOPA) (Sulzer and Zecca 2000). Because of the selective vulnerability of neuromelanin-containing neurons, dysregulation of DA homeostasis has been suggested to underlie neurodegeneration in PD (Edwards 1993; Gainetdinov et al. 1998a; Uhl 1998; Sulzer 2001; Lotharius and Brundin 2002).

#### 2 DA Metabolism and Toxicity

Accumulation of cytosolic DA (DA<sub>cvt</sub>) is toxic to cells in vitro (Mytilineou et al. 1993; Pardo et al. 1995; Sulzer et al. 2000; Fuentes et al. 2007; Mosharov et al. 2009) and several mechanisms for DA-mediated toxicity have been proposed. Spontaneous oxidation of DA in the presence of molecular oxygen produces hydrogen peroxide, hydroxyl radical, and superoxide (Graham 1978; Sulzer and Zecca 2000). Oxidation of the catechol moiety of DA yields DA-o-quinone and dopaminochrome (Sulzer and Zecca 2000); the latter can undergo a one-electron reduction to leukoaminochrome o-semiquinone radical that has been shown to produce other reactive species (Paris et al. 2009). Furthermore, DA-o-quinone reacts with free cysteine and exposed cysteine residues of proteins and glutathione, producing 5-S-cysteinyl-DA following cleavage by proteases. 5-S-cysteinyl-DA can undergo further oxidation and is toxic to cultured cells (Spencer et al. 2002) or when injected into the mouse brain (Zhang and Dryhurst 1994). 5-S-cysteinyl adducts of DA and its metabolites are readily detected in human SNpc confirming the physiological relevance of DA-induced protein damage (Fornstedt et al. 1989; Montine et al. 1995; Hastings and Berman 1999).

Other mechanisms of DA-mediated neurotoxicity include stabilization of toxic  $\alpha$ -synuclein protofibrils (Rochet et al. 2004; Mosharov et al. 2006), inhibition of chaperone-mediated autophagy by DA-synuclein adducts (Martinez-Vicente et al. 2008), and reactions of DA with nitric oxide (Daveu et al. 1997), peroxynitrite



**Fig. 1** DA homeostasis. Abbreviations: *AADC* aromatic L-amino acid decarboxylase, *AD* aldehyde dehydrogenase, *DA* dopamine (cytosolic, vesicular, and extracellular pools), *DAT* DA uptake transporter, *DA-Cys* 5-S-cysteinyl-DA, *DA-Q* DA-o-quinone, *DOPAC* 3,4-dihydroxyphenylacetate, *DOPAL* 3,4-dihydroxyphenylacetaldehyde, *L-DOPA* 3,4-dihydroxy-L-phenylalanine, *MAO* monoamine oxidase, *NM* neuromelanin inside autophagic vacuoles, *ROS* reactive oxygen species, *TH* tyrosine hydroxylase, *Tyr* tyrosine, *VMAT* vesicular monoamine transporter

(Daveu et al. 1997; Vauzour et al. 2008), and aldehydes (Collins and Bigdeli 1975; Deitrich and Erwin 1980; Naoi et al. 1993; Marchitti et al. 2007). Several recent reports confirm that a buildup of  $DA_{cyt}$  is indeed sufficient to induce progressive nigrostriatal neurodegeneration in rodents ((Caudle et al. 2007; Chen et al. 2008); also see below), although clinical studies of L-DOPA toxicity produced controversial results (Fahn et al. 2004; Holford et al. 2006; Chan et al. 2007).

Considering the toxic potential of DA, it is not surprising that multiple cellular mechanisms exist to ensure that DA concentration in the cytosol does not increase uncontrollably. It has been suggested that neuromelanin biosynthesis acts as one of the mechanisms for regulating toxic DA by-products by sequestering them into autophagic vacuoles (Sulzer and Zecca 2000). Other mechanisms include feedback inhibition of DA synthesis, catabolic DA cleavage, and synaptic vesicle sequestration.

#### 2.1 Synthesis

Catecholamine neurotransmitters DA, norepinephrine, and epinephrine are defined by the presence of a catechol ring and an amine side-chain; together with indoleamine serotonin, these compounds are referred to as monoamines. DA and other catecholamines are synthesized from the nonessential amino acid tyrosine by a series of enzymatic reactions (Fig. 1).

In the first and typically rate-limiting step, tyrosine hydroxylase (TH) attaches a hydroxyl group to the aromatic ring of tyrosine using molecular oxygen, tetrahydropterin, and ferrous iron  $(Fe^{2+})$  as substrates and cofactors (Ramsey and Fitzpatrick 2000). TH activity is regulated on transcriptional, translational, and posttranslational levels (reviewed in (Goldstein and Lieberman 1992; Kumer and Vrana 1996; Fitzpatrick 2000; Daubner et al. 2011). The latter includes phosphorylation-dependent activation of TH by various kinases and its inhibition by DA, which competes for the binding site with tetrahydrobiopterin. This feedback mechanism limits DA production when its cytosolic concentration increases.

The second enzyme in DA biosynthesis, aromatic L-amino acid decarboxylase (AADC), uses pyridoxal phosphate as a cofactor and converts L-DOPA to DA. AADC activity can be regulated by second messenger systems; antagonists acting on extracellular DA receptors enhance AADC activity and agonists diminish it (reviewed in Hadjiconstantinou and Neff (2008)). Again, these mechanisms act to decrease DA production when its extracellular concentration increases. We recently found that AADC activity is increased by cytosolic Ca<sup>2+</sup> (Mosharov et al. 2009). Although under normal conditions this mechanism may help to boost DA availability during high exocytotic activity, it may also increase the vulnerability of catecholaminergic cell populations that rely on Ca<sup>2+</sup>- driven pacemaking, i.e., SNpc and LC neurons (Hetzenauer et al. 2006; Surmeier et al. 2010).

Importantly, AADC is not saturated under physiological conditions and increased L-DOPA concentration leads to its immediate conversion to DA. As L-DOPA is able to cross the blood–brain barrier, in the early 1960s it was introduced as an effective medication for PD (Hornykiewicz 2010). L-DOPA (levodopa), frequently in combination with the peripheral inhibitors of AADC carbidopa or benserazide, alleviates cardinal symptoms of PD and has remained the most commonly used drug for the treatment of the disease. L-DOPA treatment, however, does not halt the progression of PD, due to the ongoing loss of striatal terminals that store and release DA.

#### 2.2 Cleavage

DA can be catabolized in the brain in a three step reaction that yields homovanillic acid (HVA) as a final product. First, monoamine oxidase (MAO) deaminates DA to produce 3,4-dihydroxyphenylacetaldehyde (DOPAL), which is further oxidized to 3,4-dihydroxyphenylacetate (DOPAC) by aldehyde dehydrogenase. In the next reaction catalyzed by catechol-O-methyltransferase (COMT), a methyl residue is attached to one of the hydroxyl groups of the catechol ring preventing its possible auto-oxidation to quinone. COMT however is located in postsynaptic neurons and glial cells thus playing limited role in the clearance of DA in catecholaminergic neurons (Kaakkola et al. 1987; Myohanen et al. 2010).

Of the first two reactions, MAO-mediated deamination of DA that takes place on the outer membrane of the mitochondria is the rate-limiting step. There are two isoforms of the enzyme: MAO-A is expressed in catecholaminergic cells and MAO-B is more abundant in glial cells and serotonergic and histaminergic neurons (Westlund et al. 1985). Even though MAO constitutes the first step of detoxification of DA, it produces two highly reactive compounds, DOPAL and hydrogen peroxide  $(H_2O_2)$ , which can react with each other producing hydroxyl radical. Although aldehyde dehydrogenase activity is very high and maintains the steady-state concentration of DOPAL at low levels, possible neurotoxicity of the aldehyde metabolites of amines was predicted 60 years ago due to their extremely reactive nature (Blaschko 1952). Indeed, the presence of DOPAL and its conjugation products in the brain and their toxicity has been experimentally demonstrated both in vitro and in vivo (reviewed in (Burke et al. 2004)). The proposed mechanism of toxicity involves formation of free radicals followed by the opening of the mitochondrial permeability transition pore and apoptosis.

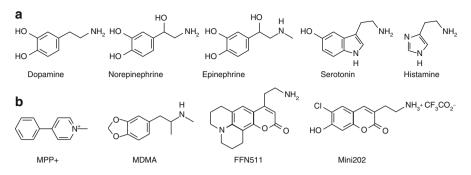
DOPAC, the final product of intracellular DA cleavage, possesses a catechol ring and, similar to DA and L-DOPA, can auto-oxidize to a quinone that further reacts with cysteine residues of proteins and glutathione. Cysteine conjugated catechols, 5-S-cysteinyl-DOPAC, 5-S-cysteinyl-DA, and 5-S-cysteinyl-DOPA, are considered to be markers of excess cytosolic DA and oxidative stress in vivo (Hastings and Berman 1999; Caudle et al. 2007).

Overall, intracellular DA catabolism may not provide neuroprotection because of the reactive nature of the DA cleavage products. Interestingly, two MAO inhibitors that cross the blood-brain barrier, selegiline and rasagiline, have been successfully tested as neuroprotective therapy in PD either as monotherapy or in combination with L-DOPA. It has been suggested, however, that the neuroprotective effect of these drugs does not depend on inhibition of MAO but rather is ascribed to the stabilization of mitochondria and the induction of antiapoptotic Bcl-2 protein family and neurotrophic factors (reviewed in Naoi and Maruyama (2010); Weinreb et al. (2010)).

#### 2.3 Synaptic Vesicle Sequestration

Monoamines are stored at high concentrations in secretory vesicles. These organelles provide protection against degradation by metabolic enzymes and enable regulated release via exocytosis. Moreover, acidic pH of the vesicles prevents auto-oxidation of DA allowing to accumulate high intravesicular neurotransmitter concentrations. Vesicular monoamine transporters (VMAT) responsible for DA uptake into synaptic vesicles are members of the mammalian major facilitator superfamily (MFS) which show structural and functional homology to bacterial toxin-extruding exporters (TEXAN) (Yelin and Schuldiner 1995; Fluman and Bibi 2009). In prokaryotes, multidrug resistance transporters are located on the plasma membrane and remove toxins from cytosol into the extracellular milieu, whereas in eukaryotes they are found on membranes of various cellular organelles and transport their substrates from cytosol into endocytic compartments.

In accordance with the toxin-extruding nature of VMAT, mammalian cDNA of the transporter was cloned by Robert Edward's laboratory on the basis of the protection that it provided against 1-methyl-4-phenylpyridinium (MPP<sup>+</sup>).



**Fig. 2** Examples of (a) native and (b) synthetic substrates of VMAT. Abbreviations:  $MPP^+$  1methyl-4-phenylpyridinium, MDMA 3,4-Methylenedioxymethamphetamine, ecstasy, FFN511false fluorescent transmitter 511, the first published optical probe designed to be a VMAT substrate, Mini202 a fluorescent ratiometric VMAT substrate sensitive to changes in vesicular pH

This neurotoxin is an active metabolite of 1-methyl-4-phenyl-1,2,3,6tetrahydropyridine (MPTP), a compound that induces parkinsonism in humans and is used in animal models of PD. VMAT-mediated resistance results from the removal of the neurotoxin from cytosol into vesicular organelles and is blocked by VMAT inhibitor reserpine (Liu et al. 1992a, b).

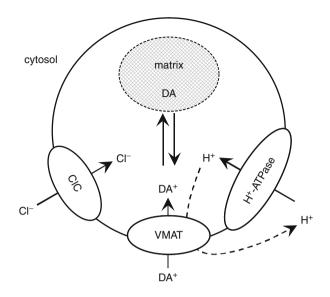
There are two isoforms of mammalian VMAT that are encoded by two separate genes that share 60 % homology (reviewed in (Guillot and Miller 2009; Wimalasena 2011)). The more abundantly expressed VMAT2 is found in central and sympathetic neurons, mast cells, and histamine-releasing cells in the gut, whereas VMAT1 expression is limited to the large dense core vesicles of peripheral endocrine cells, including adrenal chromaffin cells where it is co-expressed with VMAT2 (Weihe et al. 1994; Peter et al. 1995; Erickson et al. 1996).

Substrate specificity of VMATs is surprisingly poor, and in addition to native transmitters, it can transport a diverse range of synthetic substrates (Fig. 2; see (Wimalasena 2011) for review). It appears that two principle structural constrains that define whether a substance is a VMAT substrate are the presence of the aminoethyl group and of an aromatic ring. Such substrate promiscuity was recently exploited to develop a new set of fluorescent VMAT substrates, FFN511 (Gubernator et al. 2009) and Mini202 (Lee et al. 2010), that provide optical tools for studies of stimulation-dependent exocytosis in catecholaminergic cells (Fig. 2).

VMAT-mediated translocation of DA across secretory vesicles membrane depends on the electrochemical gradient (Johnson 1988):

$$\operatorname{Log}\left(\frac{\mathrm{DA}_{\operatorname{ves}}}{\mathrm{DA}_{\operatorname{cyt}}}\right) = \frac{\Delta \Psi \cdot F}{R \cdot T} + 2 \cdot \Delta \mathrm{pH},$$

where  $DA_{ves}$  and  $DA_{cyt}$  are catecholamine concentrations inside and outside the vesicle,  $\Delta \Psi$  is electrical gradient, *F* is Faraday's constant, *R* is the universal gas



**Fig. 3** VMAT-mediated uptake of DA into synaptic vesicles. VMAT exchanges two protons that move from vesicle to the cytosol for one positively charged DA molecule that moves from cytosol into the vesicle. Steep intravesicular proton gradient is generated by V-type H<sup>+</sup>-ATPase that uses the energy of ATP hydrolysis to achieve vesicular pH of ~5.5. Chloride channels (*ClC*) help to dissipate positive charge resulting from the accumulation of protons and catecholamines inside the vesicles. Similarly, vesicular uptake of negatively charged glutamate in some midbrain DA neurons that co-express VMAT2 and VGLUT2 increases the amount of stored DA. As small synaptic vesicles have diameters of ~50 nm and release 1,000–3,000 molecules of DA during exocytosis (Pothos et al. 1998), vesicular concentration of neurotransmitter should reach 25–75 mM. Interestingly, in such a small organelle, pH 5.5 corresponds to 0.1 protons per vesicle

constant, *T* is absolute temperature, and  $\Delta pH$  is the difference in pH between the cytosol and the vesicle.

Both the electrical and the chemical parts of the gradient are important for the transport (reviewed in (Edwards 2007; Guillot and Miller 2009)). The chemical gradient of protons is created by the vesicular proton pump, V-type H<sup>+</sup>-ATPase (Fig. 3), and VMAT uses this energy by exchanging two protons for one DA molecule. As DA is positively charged, the electrical gradient also becomes important. The influx of Cl<sup>-</sup> via the chloride channels neutralizes the buildup of positive charges due to the accumulation of DA and protons inside the vesicles therefore promoting neurotransmitter uptake. Work on isolated vesicles demonstrated that vesicle acidification is indeed much more efficient in the presence of chloride ions (Johnson 1988). Similarly, it has been shown that the knockout of the ClC-3 channel expressed on the vesicular membrane impairs vesicle acidification and induces degeneration of neuronal terminals in vivo (Stobrawa et al. 2001; Jentsch et al. 2005).

Recent data (Hnasko et al. 2010; Stuber et al. 2010) indicate that a subpopulation of midbrain neurons that project to the nucleus accumbens shell co-express VMAT2 and vesicular glutamate transporter 2 (VGLUT2). Vesicular uptake of negatively charged glutamate was found to promote accumulation of DA in the same vesicles, probably by dissipating  $\Delta \Psi$  and increasing the pH gradient that drives VMAT2-mediated monoamine transport. This suggests yet another presynaptic mechanism that regulates dopaminergic transmission.

At a steady state, when  $\Delta \Psi$  and pH in the cytosol are constant, a linear correlation is expected (i) between  $DA_{ves}$  and  $DA_{cyt}$  and (ii) between  $DA_{ves}$  and the square of the proton concentration in the vesicle:

$$DA_{ves} = A \cdot DA_{cyt} \cdot [H^+]_{ves}^2$$
, where  $A = 10^{\frac{\Delta \Psi \cdot F}{R \cdot T}} / [H^+]_{cvt}^2$ 

Indeed, both relationships have been confirmed using optical measurements of vesicular pH and electrochemical detection of  $DA_{cyt}$  and the amount of the transmitter released during stimulation-dependent vesicle exocytosis (Pothos et al. 1998; Markov et al. 2008; Mosharov et al. 2009). It is important to note, however, that the total amount of stored DA can be much higher than the amount calculated from  $DA_{ves}$  because of the presence of the vesicular "matrix" (Fig. 3). In large dense core vesicles that store norepinephrine and epinephrine, negatively charged chromogranins and secretogranins bind catecholamines allowing up to 0.5 M concentrations of vesicular transmitter (Albillos et al. 1997), which would be impossible if the osmotic and the electrical gradient were not dissipated. It remains an open question whether similar polyelectrolyte proteins play a role in DA storage in small synaptic vesicles, although the high concentrations of vesicular transmitter seem to imply this (Fig. 3).

#### 2.4 Reuptake

DA released by stimulation-dependent exocytosis acts on pre- and postsynaptic DA receptors, and its action is terminated by (1) passive diffusion, (2) COMT-mediated catabolism, and (3) reuptake by the dopamine uptake transporter (DAT).

DAT-mediated reuptake of DA has received considerable attention because this transporter is targeted by psychostimulants and drugs of abuse (reviewed in (Schmitt and Reith 1187; Zhu and Reith 2008; Sulzer 2011)). DAT activity and surface expression is regulated by phosphorylation/dephosphorylation in response to changes in extracellular DA concentration. Reuptake of DA and termination of its action on extracellular receptors also plays an important role in regulating intracellular DA pools, as illustrated in DAT-deficient mice (Gainetdinov et al. 1998b; Jones et al. 1998). In addition to expected significant elevation of extracellular DA levels, the tissue content of dopamine in DAT knockout striatum was less than 5 % of wild-type levels. It was suggested that recycled DA contributes

significantly to the maintenance of vesicular stores of DA in striatal projections of midbrain neurons. Alternatively, the observed reduction of DA intracellular stores may be due to hyperactivation of DA autoreceptors and the consequent inhibition of DA synthesis discussed above.

The importance of balanced maintenance of DA concentration in neuronal cytosol was demonstrated in a transgenic mouse that was engineered to express DAT on the plasma membrane of glutamatergic medium spiny neurons (MSNs) (Chen et al. 2008). MSNs constitute the majority of the neuronal population in the striatum and normally are exposed to DA only from the outside. In these transgenic animals however, MSNs can take up DA released by striatal projections of the midbrain dopaminergic neurons. Within several weeks following the induction of DAT expression, the mice develop motor dysfunctions, significant striatal and cortical atrophy, and progressive neurodegeneration of MSNs accompanied by oxidative stress, protein damage, and decreased glutathione levels. L-DOPA treatment worsened these symptoms, while reducing DA input to the striatum by lesioning the medial forebrain bundle had the opposite effect, demonstrating the dependence of the pathology on DA supply.

#### 2.5 Spatial Considerations

The contribution of vesicular DA uptake differs significantly between the cell bodies and axonal terminals of neurons as the later have smaller cytosolic volume and higher concentration of vesicles. Because of this difference in stored-to-total DA ratio,  $DA_{cyt}$  in synaptic terminals will be more dependent on VMAT-mediated DA uptake, which can be illustrated by a simple calculus. If one synaptic vesicle releases all its content inside the neuronal soma, the change in  $DA_{cyt}$  will equal  $DA_{ves}/N_A/V_{cell}$ , where  $DA_{ves}$ , vesicular quantal size, is ~2,000 molecules (Pothos et al. 1998);  $N_A$  is Avogadro's Constant (6.023\*10<sup>23</sup> molecules/mole); and  $V_{cell}$  is the cell soma volume (~1.77 pL for a cell with a diameter of 15 µm). It follows that emptying one vesicle would increase  $DA_{cyt}$  by ~2 nM. If, however, vesicular content is emptied inside a synaptic terminal with the diameter of ~1 µm,  $DA_{cyt}$  will be increased by >6 µM: a 3,000-fold difference.

We have recently investigated the contribution of DA metabolic pathways in the maintenance of cytosolic neurotransmitter concentration in cultured chromaffin cells (Mosharov et al. 2003) and neuronal cell bodies (Mosharov et al. 2009). Adrenal chromaffin cells, which lack neurites and are filled with secretory vesicles that each contain ~1,000,000 epinephrine or norepinephrine molecules (Wightman et al. 1991), are closer to synaptic terminals in their stored-to-total DA ratio. Cytosolic catechol-amine concentration in these cells is 10–20  $\mu$ M and can be increased severalfold by L-DOPA treatment. Consistent with strict control of the level of cytosolic catechol-amine concentration, inhibition of either MAO or VMAT does not alter cytosolic transmitter levels, presumably because one pathway compensates for decreased activity of the other. When, however, both MAO and VMAT are blocked simultaneously, cytosolic catecholamine levels increase by more than sixfold.

In contrast to chromaffin cells,  $DA_{cyt}$  in neuronal cell bodies is below the limits of detection (<0.1  $\mu$ M) unless augmented by L-DOPA pretreatment (Mosharov et al. 2009). Importantly,  $DA_{cyt}$  increases readily upon treatment with MAO inhibitors, but no contribution of somatic VMAT-mediated uptake can be detected either in the absence or in the presence of MAO blockade. Obviously, some DA secretory organelles reside in or close to neuronal cell bodies, as evident from somatodendritic DA release from midbrain neurons (Rice et al. 1997; Jaffe et al. 1998); however, the number of DA storage vesicles in the soma is probably insufficient to have significant effect on  $DA_{cyt}$  following VMAT blockade.

Amphetamine and its derivatives are often used to disrupt vesicular and DA<sub>cvt</sub> pools. These psychostimulants have multiple intracellular sites of action, including inhibition of MAO, stimulation of catecholamine synthesis, redistribution of catecholamines from secretory vesicles to the cytosol, and induction of catecholamine reverse transport from the cytosol to the extracellular milieu via dopamine and norepinephrine uptake transporters (reviewed in (Sulzer 2011)). Chromaffin cells acutely treated with methamphetamine (METH) display decreased quantal size (Sulzer et al. 1995; Anderson et al. 1998; Mundorf et al. 1999) and increased cytosolic catecholamine levels, indicating that effects of METH on DA metabolism and the leakage of DA from vesicles are the dominant processes in these cells (Mosharov et al. 2003). In contrast, neurons exposed to METH showed decreased DA<sub>cvt</sub> in the somas (Mosharov et al. 2009), demonstrating that DAT-mediated reverse transport governs the steady-state DA concentration, while redistribution of neurotransmitter from vesicles has negligible effect in neuronal cell bodies. While it is currently impossible to measure cytosolic DA in axonal terminals due to their small size, their similarity to chromaffin cells suggests a possibility that an equally large increase in DA<sub>cvt</sub> occurs in METH-treated neuronal terminals. If so, this may explain a significant increase in striatal 5-S-cysteinyl-DA levels (LaVoie and Hastings 1999) and the pattern of METH-induced neurodegeneration in which neurites but not cell bodies are damaged in vitro (Cubells et al. 1994; Larsen et al. 2002), in animal models in vivo (Ricaurte et al. 1982), and in imaging studies of METH abusers after prolonged abstinence (Sang et al. 2007).

#### 2.6 Duration of Exposure

The time during which a cell is exposed to the toxic effect of DA and its metabolites is another parameter that significantly affects neurotoxicity. In vitro, DA-mediated neurotoxicity depends linearly on the dose (concentration\*time) of cell exposure to elevated  $DA_{cyt}$  levels, i.e., neurons can be quite resistant to short bursts of high  $DA_{cyt}$  but are compromised if small elevations in cytosolic transmitter persist for sufficiently long time (Mosharov et al. 2009). This consideration becomes very important when comparing data obtained from experiments in cellular and animal models with neuropathology in human as even the early onset type of PD develops over decades of life, which is orders of magnitude longer than the life span of rodents and cells in culture.

#### **3 VMAT Expression and DA-Mediated Neurotoxicity**

#### 3.1 Less VMAT, More Toxicity

Considerable experimental evidence supports the idea that decreased vesicular uptake correlates with enhanced toxicity. Reserpine, an inhibitor of VMAT, induces parkinsonism by depleting releasable pool of DA from striatal terminals (Carlsson et al. 1957). The striatum of animals treated with reserpine shows marked rise in 5-S-cysteinyl-DA and oxidized glutathione levels, indicative of elevated cytosolic DA production and an ongoing oxidation stress (Fornstedt and Carlsson 1989; Spina and Cohen 1989; Bilska et al. 2007). Although no dopaminergic cell loss has been reported in reserpinized animals, a 7-day treatment with tetrabenazine, another VMAT inhibitor, induces irreversible histological changes in the neurons of SNpc (Satou et al. 2001). In the amphetamine-induced model of axonal degeneration in the striatum, reserpine exacerbated drug-induced toxicity, whereas pretreatment with TH inhibitor  $\alpha$ -methyl-p-tyrosine provided protection (Wagner et al. 1983). Additionally, reserpine causes concentration-dependent death of RCSN-3 cells, a catecholaminergic cell line derived from adult rat SNpc (Fuentes et al. 2007).

Several genetic mouse models with various levels of VMAT2 expression have been created to study the role of the transporter in neurotoxicity (see (Guillot and Miller 2009; Taylor et al. 2011) for comprehensive reviews). Mice with a homozygous deletion of VMAT2 transporter (VMAT $2^{-/-}$ ) do not show any gross anatomical abnormalities, including normal number of neurons in SNpc, but die within a few days after birth, probably because they do not feed (Fon et al. 1997; Takahashi et al. 1997; Wang et al. 1997). The brains of VMAT2<sup>-/-</sup> animals manifest severely compromised DA storage ( $\sim 1.5$  % of wild-type levels). Midbrain neuronal cultures from these mice show normal morphology and TH fibers density but contain only 3 % of the intracellular DA levels of the VMAT2<sup>+/+</sup> cultures and are unable to accumulate DA within synaptic vesicles (Fon et al. 1997; Yamamoto et al. 2007). Importantly, VMAT2<sup>-/-</sup> neurons are significantly more sensitive to METH-induced neurodegeneration of primarily neurites and display intense labeling for ROS formation upon drug treatment (Larsen et al. 2002). These data indicate that although METH-induced oxidative stress does not require the presence of a vesicular pool of DA, vesicular sequestration of the neurotransmitter provides significant neuroprotection.

In contrast to animals with complete VMAT2 deletion, VMAT2 heterozygous mice (VMAT2<sup>+/-</sup>) are viable to adulthood. VMAT2<sup>+/-</sup> mice display  $\sim 50 \%$  reduced total DA, diminished levels of extracellular striatal DA, as well as decreased potassium- and amphetamine-evoked DA release in vivo and in postnatally cultured cells (Fon et al. 1997; Takahashi et al. 1997; Wang et al. 1997). Although the animals perform normally on motor tests, they show depressive-like phenotype and impairments in learned helplessness and conditioned place preference paradigms (Takahashi et al. 1997; Wang et al. 1997; Fukui et al. 2007). As expected, mice are more sensitive to neurodegeneration induced by

methamphetamine and MPTP (Gainetdinov et al. 1998a; Fumagalli et al. 1999), confirming the ability of VMAT2 to protect cells by removing toxins from the cytosol. Though no PD-like neuropathology was observed in VMAT2<sup>+/-</sup> mice up to 7-month of age, further studies are required to determine whether neurodegeneration of the nigrostriatal system may occur later in life. Interestingly, while no loss of the nigrostriatal biochemical markers was found in VMAT2<sup>+/-</sup> animals injected with L-DOPA for 28 days (Reveron et al. 2002), cultured DA neurons from these mice show increased vulnerability to the toxic effect of L-DOPA (Kariya et al. 2005).

Another line of mice that proved extremely helpful in PD research expresses only 5 % of the normal level of VMAT2. The first generation of mice that have a hypomorphic allele of the VMAT2 (VMAT2-deficient KA1 line (Mooslehner et al. 2001)) was later found to have a spontaneous deletion of the alphasynuclein gene (Specht and Schoepfer 2001; Colebrooke et al. 2006). As this protein plays an important role in the pathology of PD (reviewed in (Vila and Przedborski 2004; Eriksen et al. 2005; Sulzer 2007; Dawson et al. 2010)), further breeding was done to remove all traces of alpha-synuclein deficiency.

The resulting VMAT2-LO mice show 95 % reduction of VMAT2 expression and 85 % reduction in DA levels in the striatum (Caudle et al. 2007). As the animals age, they show signs of PD-like progressive neurodegeneration, including L-DOPA-responsive motor deficits; oxidative stress and protein damage; decreased DA, DAT, and TH levels in the striatum; and pathological accumulations of alphasynuclein and a reduced number of DA neurons in the SNpc (Caudle et al. 2007; Taylor et al. 2011). Importantly, animals do not manifest motor symptoms and some of the biochemical markers until they are 12–28-month-old (Taylor et al. 2011), highlighting the slow nature of progressive neurodegeneration in this animal model.

The onset of the motor symptoms in PD is accompanied and generally preceded by a variety of nonmotor disturbances, including but not limited to olfactory deficits, constipation, sleep disorders, and depression. It has been suggested that degeneration of other monoamine systems, such as norepinephrine neurons in LC and serotonin neurons in the dorsal raphe nucleus, may contribute to the development of the nonmotor symptoms (reviewed in (Dauer and Przedborski 2003; Fornai et al. 2007; Rommelfanger and Weinshenker 2007; Fahn 2008; Obeso et al. 2008; Delaville et al. 2011; Lang 2011)). Biochemical characterization of VMAT2-LO mice revealed progressive depletion of all monoamines (Caudle et al. 2007; Taylor et al. 2009) as well as age-dependent neurodegeneration in the LC that preceded the loss of neurons in the SNpc (Taylor et al. 2011), similar to the clinical data from PD patients. Moreover, VMAT2-LO mice demonstrate early deficits in olfactory discrimination, delayed gastric emptying, altered sleep latency, anxiety-like behavior, and age-dependent depressive behavior (Taylor et al. 2009), confirming the involvement of monoamine dysfunction in the manifestation of nonmotor symptoms of PD. Overall, the VMAT2-LO mouse model recapitulated the majority of motor and nonmotor symptoms of PD and demonstrates that a reduced capacity of cells to sequester DA and other monoamines inside storage vesicles is sufficient to cause PD-like neurodegeneration of neurons and their axonal projections.

#### 3.2 More VMAT, Less Toxicity

As VMAT2 should prevent DA-induced toxicity by sequestering neurotransmitter inside synaptic vesicles, enhancing VMAT-mediated transport is expected to be neuroprotective. Clinical data indirectly support this claim as neurons in the ventral tegmental area, a dopaminergic cell population that is spared in PD, express higher levels of VMAT2 and produce lower amounts of neuromelanin compared to neurons in SNpc (Hirsch et al. 1988; Zecca et al. 2001; Liang et al. 2004). Furthermore, higher expression level of VMAT2 compared to that of DAT has been suggested to predict lower vulnerability of neuronal populations in PD (Uhl 1998; Miller et al. 1999), while gain-of-function haplotypes in the human VMAT2 gene reduce the risk of developing PD in women (Glatt et al. 2006).

The link between neurotoxicity and the activity of VMAT2 has been directly studied in vitro using cultured cells. As described above, mouse midbrain neurons treated for several days with DA precursor L-DOPA produce neuromelanin due to accumulation of cytosolic DA and downstream oxidation products (Sulzer et al. 2000). Virus-mediated overexpression of VMAT2 in these cultures prevents neuromelanin synthesis, increases the number of DA molecules packaged per synaptic vesicle, decreases DA<sub>cvt</sub>, and efficiently protects neurons from L-DOPAinduced cell death (Pothos et al. 2000; Sulzer and Zecca 2000; Mosharov et al. 2009). Similarly, overexpression of VMAT2 in primary rat mesencephalic cultures and PC12 cells, a pheochromocytoma-derived cell line, increases intracellular DA content, augments stimulation-dependent DA release, and attenuates cell death induced by METH; downregulation of VMAT2 using virally delivered shRNAs produces an opposite effect (Vergo et al. 2007). Additionally, increased expression of VMAT2 in vivo is able to mitigate the oxidative stress and gliosis caused by METH (Guillot et al. 2008). Together these data strongly support the inverse relationship between DA-induced neurotoxicity and the activity of VMAT2.

#### 4 Conclusion

Overview of enzymes and transporters involved in DA homeostasis illustrates that all metabolic steps work to maximize the amount of DA available for exocytosis while at the same time limiting the accumulation of potentially toxic cytosolic transmitter. Importantly, all except one metabolic step in DA turnover affect neurotoxicity and neurotransmission in the same direction, i.e., by altering the levels of cytosolic and vesicular DA in synchrony. For example, increased activities of TH and AADC will enhance DA-mediated toxicity and the amount of DA stored in vesicles, whereas increased catabolism will diminish them both. The sole exception is the activity of VMAT, which has opposing effects on cytosolic and vesicular DA levels. This unique position of VMAT in DA handling provides therapeutic advantages that can be explored in future drug development.

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# Rotenone as Preclinical Model Compound in Parkinson Disease

# Jason R. Cannon and J. Timothy Greenamyre

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#### Abstract

The motor symptoms of Parkinson's disease (PD) are characterized by a combination of bradykinesia, resting tremor, rigidity, and postural instability, which are caused by striatal dopamine depletion resulting from loss of nigral dopamine neurons and dopaminergic projections. At the time of diagnosis,

J.R. Cannon (🖂)

J.T. Greenamyre The Pittsburgh Institute for Neurodegenerative Diseases, University of Pittsburgh, Pittsburgh, PA, USA e-mail: jgreena@pitt.edu

School of Health Sciences, Purdue University, West Lafayette, IN, USA e-mail: cannonjr@purdue.edu

there is already extensive loss of striatal dopamine and nigral dopamine neurons. Patients also commonly experience a host of cognitive, psychiatric, autonomic, sleep, and sensory disturbances that likely result from pathology to regions outside the substantia nigra. These nonmotor abnormalities may occur decades before diagnosis. "Preclinical" PD is the disease before motor symptom severity elicits a PD diagnosis. Currently, there is an emphasis on the creation and characterization of preclinical models to identify early-stage PD features and test early intervention strategies. The rat rotenone model tests the hypothesis that nigral dopamine neurons are selectively sensitive to systemic mitochondrial complex I inhibition. The model replicates selective loss of nigral dopamine neurons and was the first neurotoxicant model to reproduce intracellular  $\alpha$ synuclein accumulation, similar to Lewy bodies. Subsequent investigations have found the model to replicate many additional PD features. In this chapter, the utility of this model to examine preclinical PD endpoints is evaluated. The major emphasis is pathological and functional alterations that are known to occur in the earliest stages of PD or even be predictive of disease development. Finally, specific rotenone administration regimens to produce preclinical PD models are detailed. In summary, available data suggest that the rat rotenone model is an excellent system to examine preclinical PD.

#### **Keywords**

Parkinson's disease • Preclinical • Rotenon	e
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List of Abb	reviations
6-OHDA	6-hydroxydopamine
AD	Alzheimer's disease
DNA	Deoxyribonucleic acid
LBD	Lewy body dementia
MIBG	Metaiodobenzylguanidine
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine
PD	Parkinson's disease
PET	Positron emission tomography
RBD	REM behavior disorder
REM	Rapid eye movement
SPECT	Single-photon emission computed tomography

## 1 Introduction

Parkinson's disease (PD) is characterized behaviorally by the presence of a combination of the cardinal motor features that include bradykinesia, resting tremor, rigidity, and postural instability. These key motor deficits are caused by striatal dopamine depletion that results from the loss of nigral dopamine neurons and their striatal dopaminergic projections (Bertler et al. 1958; Carlsson et al. 1958;

Moore et al. 1971). A major pathological hallmark is the presence of cytoplasmic inclusions known as Lewy bodies in surviving dopamine neurons (Forno 1996). Aggregated  $\alpha$ -synuclein and ubiquitin are major components of Lewy bodies, and mutations in  $\alpha$ -synuclein that promote aggregation or in proteins that affect proteasomal activity cause PD (Morris 2005; Spillantini et al. 1997). In addition to pathology in the nigrostriatal dopamine system, pathology is evident in other brain regions, particularly in brainstem nuclei, where  $\alpha$ -synuclein accumulation occurs and is thought to temporally precede nigral aggregation (Braak et al. 2003; Braak et al. 2004). Systemic pathology also occurs, especially in the enteric nervous system (Braak et al. 2006; Wakabayashi et al. 1988). PD patients also commonly experience a host of cognitive, psychiatric, autonomic, sleep, and sensory disorders that likely result from pathology to neuronal regions outside the substantia nigra (Adler 2005; Emre 2004; McDonald et al. 2003; Papapetropoulos and Mash 2005).

Pharmacologic and surgical approaches can effectively treat many of the motor symptoms. However, current treatment options do not slow or prevent disease progression. PD is currently thought to affect at least 5 million individuals worldwide (Dorsey et al. 2007). The most significant risk factor for PD is ageing (Collier et al. 2011). Thus, given that the average life span is rising, in part due to medical advances related to cardiovascular diseases and cancers, PD incidence is expected to increase in an ageing population.

The causes of most cases of PD remain unknown. Genetic factors continue to be identified. However, monogenetic causes likely account for less than 10 % of total PD cases. Environmental exposures have long been linked to PD, with numerous structural and use classes of compounds linked to PD and other neurodegenerative diseases. While many reports have suggested links between environmental exposures and PD, identifying specific causative risk factors has been very difficult (Cannon and Greenamyre 2011). More recently specific toxicant exposures have been identified as risk factors using specialized epidemiological techniques. Notably, exposures to the herbicide paraquat and the pesticide rotenone have been identified as PD risk factors (Tanner et al. 2011). The identification of these toxicants as risk factors is supported by the fact that both of these toxicants have been utilized extensively to model PD in animals. Unfortunately, for most PD cases there is limited understanding of etiology. The majority of cases are thought to arise from a combination of genetic predisposition and environmental exposures. There is some evidence for a role of specific interactions in PD risk; however, our understanding of these interactions remains rudimentary (Cannon and Greenamyre 2012).

Thus, there are two key challenges that PD researchers and clinicians continue to have: (1) identifying causative factors (a reduction in exposure to causative factors could, in principle, reduce PD incidence over time) and (2) the development of disease-modifying therapies (such therapies would slow or halt progression in addition to treating the primary motor symptoms). While work to accomplish these goals typically involves human studies, animal models are extensively utilized during the earliest stages and throughout the process. Unfortunately, translating positive results from neurotoxicant-based animal studies to clinical trials has been largely unsuccessful. There are several possible explanations: (1) Neuroprotective

regimens are usually administered prior to toxicant insult. This type of regimen bears little relevance to a clinical setting, where a patient would be treated after diagnosis. Thus, positive studies from pretreatment regimens may have little relevance and translational impact in humans. (2) Relevance of the model. Most neurotoxicant models do an outstanding job of lesioning the nigrostriatal dopamine system. However, a model that does so in hours or a few days does not reproduce the chronic development of pathology in human PD. Further, regimens that simply prevent the primary mechanism of action of the insult in a neurotoxicant model are unlikely to be meaningful, for example, showing that an antioxidant is protective in an acute oxidative stress model (6-hydroxydopamine, 6-OHDA) or that a drug which alters 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) metabolism or metabolite accumulation in dopamine neurons is unlikely to provide substantially useful information on whether it is a viable neuroprotective agent (Cannon and Greenamyre 2010). (3) Failure to appreciate early-stage features of PD and PD models. The earlier a neuroprotective intervention is administered in the disease process, the more likely it will have a disease-modifying impact. The use of preclinical animal models can help to test potential therapeutics for an early-stage administration both directly and indirectly, by aiding in the development of early-stage diagnosis and testing whether therapeutic regimens prevent development of preclinical endpoints or are neuroprotective. Currently, preclinical animal models are limited - most models focus on reproducing the end-stage phenotype, primarily major striatal dopamine depletion and nigral dopamine cell loss. In this brief chapter, the development and use of the rotenone model to reproduce and identify preclinical features of PD is critically evaluated.

### 2 The Rotenone Model of Parkinson's Disease

Rotenone is a compound that occurs naturally in the roots of a variety of plant species. It is a highly selective inhibitor of mitochondrial complex I that has been historically used as an insecticide and to kill fish (Ravanel et al. 1984). Systemic complex I inhibition occurs in PD patients (Parker et al. 1989; Schapira et al. 1989). Thus, it has been hypothesized that toxicants that produce systemic complex I inhibition may have a role in PD. It should be noted that the classic dopaminergic neurotoxicant, MPTP, also selectively inhibits complex I. However, due to its affinity for the dopamine transporter, it selectively affects catecholaminergic neurons (Javitch et al. 1985; Nicklas et al. 1985). Conversely, rotenone is a highly lipophilic molecule able to enter all cells and produce complex I inhibition.

There have been many attempts to produce a rotenone model of PD, including high-dose systemic administration and intracerebral infusions that produce nonspecific lesions (Ferrante et al. 1997; Heikkila et al. 1985; Rojas et al. 2009). A major advancement came when systemic rotenone was chronically administered to rats at much lower doses, producing complex I inhibition similar to that observed in PD patients. This regimen produced highly selective nigral dopamine neuron degeneration (Betarbet et al. 2000). Here, for the first time in an animal model,

cytoplasmic,  $\alpha$ -synuclein-positive inclusions that have striking similarities to Lewy bodies were observed in surviving dopamine neurons. Reproduction of this key pathological feature represented a major advantage over other available neurotoxicant models. While variability in the number of animals that exhibited a lesion and variability in the magnitude and morphology of the lesion were major obstacles for using the model in neuroprotection and potentiation studies, a more recent version is much more reproducible (Cannon et al. 2009). While the rat is the most common species that rotenone has been used to model PD, it has been used in other major biological models including Drosophila, C. elegans, and zebrafish (Cannon and Greenamyre 2010). Not only has the rotenone model replicated key PD features, it has been found to predict previously unknown features of human PD. Notably, rotenone treatment in the rat produces iron accumulation in the substantia nigra through a previously unknown mechanism involving transferrin and transferrin receptor 2 (Mastroberardino et al. 2009). This study also found the same alterations in the brains of PD patients. In addition to replicating these key pathological features, the rotenone model has been found to be useful in reproducing several key preclinical features of PD, which are discussed in this chapter.

# 3 Preclinical Parkinson's Disease and the Relevance of the Rotenone Model

What is preclinical PD? Simply stated, "preclinical" is the disease before characteristic motor symptoms become severe enough that the patient enters a clinic and is diagnosed with PD. Certainly, there is much patient-to-patient variability in the threshold of symptoms that are required to reach "clinical" PD. However, it is worth noting that by the time a patient is diagnosed with PD,  $\sim$ 70–80 % of striatal dopamine and  $\sim$ 50–60 % of nigral dopamine neurons are lost (Bernheimer et al. 1973). Thus, it is thought that damage and cell loss may occur over several decades before the pathology becomes severe enough to elicit noticeable symptoms.

### 3.1 Autonomic Dysfunction

Autonomic dysfunction may precede motor deficits by decades. For example, gastrointestinal dysfunction in the form of reduced bowel movements is a risk factor for future development of PD, an average of 12 years into follow-up (Abbott et al. 2001). A second study confirmed that constipation is a risk for PD (Savica et al. 2009). Both studies conclude that gastrointestinal dysfunction can precede the PD diagnosis by 10–20 years. Interestingly, Lewy bodies and Lewy neurites are also found in the myenteric and submucosal plexuses of the gut (Braak et al. 2006; Lebouvier et al. 2008; Wakabayashi et al. 1990). While these histological examinations were performed postmortem, the temporal development of symptoms would suggest that pathology in the gut is an early event in PD.

In rats treated with low-dose rotenone, which does not elicit overt nigrostriatal toxicity (2.0 mg/kg/day, 5x/week for 6 weeks), gastrointestinal dysfunction similar to that in PD patients is observed (Drolet et al. 2009). Insoluble  $\alpha$ -synucleinpositive aggregates strikingly similar to that which occur in the gut of PD patients were found in the small intestine myenteric plexus 6 months after treatment. Neuronal loss in the gut and increased gastric transit time were also found in this study, also 6 months after rotenone treatment. Thus, chronic, low-dose systemic rotenone produces gastrointestinal dysfunction, in the absence of overt motor deficits or brain pathology. A second rat study also showed the development of delayed gastric emptying and enteric neuron dysfunction after rotenone treatment (Greene et al. 2009). Direct gastric administration of rotenone in mice is also able to produce  $\alpha$ -synuclein aggregation (Pan-Montojo et al. 2010). The pathological findings in this study also showed that pathology in the gut preceded that in the substantia nigra. Thus, chronic rotenone treatment in rodents is able to reproduce this key preclinical event that occurs in many PD patients. Beyond simply validating the ability of this model to replicate key PD features, it is expected that gastrointestinal endpoints in future studies will provide much useful information on the pathophysiology of gut abnormalities in PD and a useful tool for examining treatments relevant to autonomic symptoms.

# 3.2 Olfactory Pathology and Deficits

The olfactory system is also affected in the earliest stages of PD.  $\alpha$ -Synuclein accumulation in the anterior nucleus of the olfactory bulb precedes nigral pathology (Braak et al. 2003). Further, at onset, olfaction is typically severely impaired in the majority of PD patients, with normal function or mild impairment in other parkinsonian disorders (Hawkes 2003). Olfactory deficits may also be an important marker for Lewy body dementia (LBD), and although Alzheimer's disease (AD) patients may exhibit olfactory loss, it is typically less severe than in LBD (Olichney et al. 2005; Postuma et al. 2010). While numerous studies have found olfactory deficits to be predictive, the time where the predictive power is significant is typically 2–4 years prior to diagnosis (Postuma et al. 2010). Specificity may also be low; while those in the lowest quartile of olfactory function in the Honolulu-Asia Ageing Study had an odds ratio of 5.2 for developing PD, only 2 % of those in this quartile developed PD (Postuma et al. 2010; Ross et al. 2008).

In the rotenone model, examination of oxidative damage in the brain found that the most affected areas were the midbrain and olfactory bulb (Sherer et al. 2003). While this data was obtained from animals exhibiting severe parkinsonism, the data suggest that this brain region is a target and should be evaluated in earlier stages of the model or under low-dose regimens. Specifically, functional (olfactory thresholds and discrimination) and detailed neuroanatomical (examination of specific cell populations) studies are needed to determine how closely effects on the olfactory system resemble that which occur in human PD.

### 3.3 Pathology in Brainstem Nuclei

Pathology in the brainstem occurs in PD and is thought to precede damage to nigral dopamine neurons. In particular,  $\alpha$ -synuclein accumulation in the dorsal motor nucleus of the vagus (acetylcholinergic) and locus coeruleus (noradrenergic) occurs in the earliest stages of PD (Braak et al. 2003, 2004). Pathophysiology in these nuclei may underlie some of the nondopaminergic PD symptoms.

To date one group has reported overt neuronal loss in the locus coeruleus after systemic subcutaneous rotenone administration (Lin et al. 2008). This finding was observed along with the typical nigrostriatal dopamine neuron degeneration. A recent in vitro study found that rotenone treatment promotes  $\alpha$ -synuclein aggregation in primary neuronal cultures from the locus coeruleus (Chaves et al. 2010). A more thorough examination is needed to determine how early the locus coeruleus is affected in the rotenone model.

Mice receiving gastric rotenone administration exhibit inflammation in the dorsal motor nucleus of the vagus and  $\alpha$ -synuclein-positive aggregate formation in acetylcholine positive neurons (Pan-Montojo et al. 2010). Notably, in this time-course study, brainstem pathology occurred prior to nigral pathology. Thus, chronic rotenone treatment is able to replicate the temporal development of key preclinical brainstem pathology.

### 3.4 Cardiac Function

Cardiac function is clearly altered in PD patients. In particular, postganglionic sympathetic innervation is affected in PD and DLB patients (Postuma et al. 2010). Although it requires cardiac metaiodobenzylguanidine (MIBG) scintigraphy to measure, which is costly, PD patients typically exhibit abnormal MIBG scintigraphy in the earliest disease stages (Spiegel et al. 2007).

Rotenone-treated rats that exhibit nigrostriatal dopamine lesions have decreased baroreflex sensitivity similar to that which occurs in PD patients (Yu et al. 2008). In this study, cardiac function was assessed at the end of rotenone administration. Given that in human PD patients, attenuated baroreflex function may precede motor symptoms, it is possible that rotenone treatment may recapitulate this preclinical PD symptom (Goldstein 2011; Goldstein et al. 2007). Evaluation of cardiac function in rotenone-treated rats, before nigrostriatal lesions develop or at subthreshold doses, will help to determine if this is a preclinical feature of the model.

### 3.5 Sleep Disorders

Sleep disorders are also common in PD patients and may precede motor deficits. Most notably is REM behavior disorder (RBD), where patients do not exhibit the normal atonia that occurs in REM sleep. The result is ultimately "acting out" of dreams, which may lead to crying out, kicking, or thrashing (Gagnon et al. 2006). RBD occurs in roughly one third of PD patients, with about another one third exhibiting asymptomatic loss of atonia (Gagnon et al. 2002). RBD is also common in other neurodegenerative disorders. It is thought to stem from damage to pontine nuclei, which are affected in the early stages of PD. Further, those with RBD have a high risk for developing PD, with 40–60 % developing a neurodegenerative disorder (1/2 develop PD) 10 years after follow-up (Postuma et al. 2010). Thus, the long-lag period indicates that RBD/reduced REM atonia is a key preclinical feature of many PD cases. PD patients also exhibit insomnia and excessive daytime somnolence, also thought to arise in part due to degeneration of pontine nuclei (Braak et al. 2003; Chaudhuri et al. 2006; Postuma et al. 2010).

A few sleep-related studies have been conducted on rotenone-treated rats. The first report found that both vehicle (dimethyl sulfoxide and polyethylene glycol; 1:1) and rotenone-treated animals exhibited sleep disturbances, rendering interpretation of the direct effect of rotenone on sleep architecture difficult to discern (Garcia-Garcia et al. 2005). A later study in rotenone-treated rats found that slow-wave sleep was increased during the active time (dark) and decreased during the inactive period (light) (Yi et al. 2007). Further, REM sleep was enhanced during the dark period. Interestingly, this study found that alterations to sleep architecture were not mediated by dopamine. More detailed studies are needed to assess the temporal development of sleep disturbances in rotenone-treated rats. Muscle activity will also need to be assessed to determine if atonia occurs during REM, following rotenone treatment.

### 3.6 Oxidative Stress and Damage to Biomolecules

Oxidative stress has long been known to occur in human PD. Proving a direct causative role for oxidative stress in human PD is difficult because brain samples are obtained at autopsy. It should be noted that many of the prominent models of PD (6-OHDA, MPTP, and rotenone) utilize oxidative stress (through different mechanisms of action) to effectively lesion the nigrostriatal dopamine system and replicate the key pathogenic features of clinical PD (Betarbet et al. 2006; Przedborski and Vila 2001; Schwarting and Huston 1996). Such data would suggest that oxidative stress is important in the pathogenesis of the disease, not simply a by-product of end-stage pathology.

Diverse examples of oxidative damage in the human PD brain include increases in total iron content in the substantia nigra (thought to be a contributor to oxidative stress), superoxide dismutase (involved in cellular response to oxidative stress), lipid peroxidation (a marker for oxidative damage to lipids), protein oxidation, and DNA oxidation versus age-matched controls (Alam et al. 1997; Dexter et al. 1989; Floor and Wetzel 1998; Saggu et al. 1989; Sofic et al. 1988). However, brain oxidative stress and damage are difficult to utilize as preclinical features in humans due to sampling availability of relevant endpoints until after autopsy.

Oxidative stress and damage in PD is not limited to the brain, with evidence in serum and systemic tissues. For example, an overall index of serum oxidant status is

increased in PD patients and the antioxidant status is decreased (Forte et al. 2004). More specifically, plasma lipid peroxide levels are increased in PD patients (Kilinc et al. 1988), indicating a general systemic oxidative damage in PD. Serum levels of superoxide dismutase and glutathione peroxidase have been found in PD patients, indicating a systemic response to oxidative stress (Kalra et al. 1992). Oxidation of plasma nucleic acid is also increased in PD patients (Abe et al. 2003; Kikuchi et al. 2002). Additionally, several cell types undergo oxidative damage in PD, including platelets (Gotz et al. 2000; Krige et al. 1992; Yoshino et al. 1992), erythrocytes (Kilinc et al. 1988; Younes-Mhenni et al. 2007), lymphocytes (Migliore et al. 2001, 2002; Petrozzi et al. 2001), and neutrophils (Gatto et al. 1996; Vitte et al. 2004).

Rotenone has been found to selectively lesion nigral dopamine neurons through complex I inhibition, resulting in oxidative stress, to which these neurons are particularly susceptible (Sherer et al. 2003). Given the number of clinical PD parameters that the rotenone model replicates, it is highly likely that oxidative stress in the earliest disease stages is important in human PD pathogenesis. In vitro studies have shown that rotenone treatment leads to progressive oxidative stress in neuroblastoma cells (Sherer et al. 2002). Further, work in organotypic slice cultures showed the development of oxidative stress along with loss of nigral dopamine neurons. Both oxidative stress and neuronal loss were prevented with antioxidant treatment (Testa et al. 2005). Later work showed that DJ-1, an important oxidative stress protein, is itself oxidatively modified in nigral neurons from both lesioned and non-lesioned rotenone-treated rats (Betarbet et al. 2006). This finding suggests that oxidative stress and damage are early events occurring prior to overt neuronal loss. Specialized redox imaging performed as a time course in zebrafish larvae also shows that oxidative stress is an early event after rotenone treatment (Horowitz et al. 2011).

Systemic studies on the temporal development of oxidative stress are limited in the rotenone model. Studies in progress in the Greenamyre lab (unpublished) show that DNA damage in the blood is profound and occurs within hours of the first rotenone administration. Studies in the future on the temporal development of damage to specific cell types are expected to be highly informative.

## 4 Practical Considerations: Executing and Utilizing the Rat Rotenone Model to Assess Preclinical Endpoints

In the rat rotenone model, there are two basic regimens to assess preclinical endpoints:

 Standard dose, with early sampling. While there are certainly lab-to-lab differences, dependent on age and strain, in our experience, administration of systemic rotenone at 3.0 mg/kg/day will typically produce debilitating motor deficits ~7–14 days after initial administration (Cannon et al. 2009). Sampling at 5 days or less, before motor deficits or nigrostriatal pathology arises, allows for the assessment of preclinical endpoints. A major advantage of this approach is that it is known that at this dose, animals will eventually develop the key motor and pathological features of end-stage PD. However, sampling after an acute administration may result in the inability to detect preclinical endpoints that develop chronically.

2. Low-dose, long-term treatments. Animals treated at rotenone doses 2.0–2.5 mg/ kg/day do not typically develop severe motor deficits or nigrostriatal pathology (Cannon et al. 2009; Drolet et al. 2009). A major advantage to this approach is that the development of long-term preclinical endpoints can be assessed. For example, after 2.0 mg/kg/day rotenone, 5 days/week for 6 weeks, rotenone treatment was discontinued and rats were found to develop gastrointestinal pathology and transit deficits 6 months later (Drolet et al. 2009). A dose of 2.5 mg/kg/day was found to elicit motor symptoms in few wild-type animals, but potentiation occurred in transgenic animals expressing disease-causing mutations in  $\alpha$ -synuclein (Cannon et al. 2013). Thus, low-dose, long-term rotenone can be used to assess the development of preclinical endpoints and potentiation (especially, gene-environment interactions). Potential disadvantages are that animals likely will not eventually develop a phenotype characteristic end-stage PD and a lengthy dosing period may be required to elicit the desired preclinical features. Rats treated at 2.0–2.5 mg/kg/day, as long as 60 days, may not develop motor deficits (Cannon et al. 2009). Thus, it may be somewhat more difficult to determine the importance of potential preclinical features in a model that does not develop the key features of end-stage pathology. Further, evaluation at up to 6 months of age can be costly and time consuming. However, it should be noted that low-dose neurotoxicant exposures and chronic development are likely more relevant to human exposures and the temporal development of the disease.

# 5 Future Directions: The Need for Better Preclinical Models and Suggested Uses

Modeling preclinical features is of major importance to advancing PD research. First understanding the key preclinical features of the disease will aid in earlier diagnosis. This will be particularly important as neuroprotective agents become available in the future. The earlier a neuroprotective agent could be administered, the more likely it would be effective. Second, a better understanding of preclinical features will increase our understanding of the pathophysiology of PD and perhaps lead to new therapeutic targets.

# 5.1 How Can the Rotenone Model be Further Validated for Modeling Preclinical Use?

While it is now over a decade old, the rotenone model continues to evolve. In its most recent form, it is much more reproducible and amenable to neuroprotection and potentiation studies (Cannon et al. 2009, 2013; Tapias et al. 2010). Current data suggest that the rotenone model can replicate many of the key preclinical features. However, much more data is needed. In particular, alterations in olfactory function,

sleep architecture, and cardiac function need to be further assessed to determine when they may occur, relative to nigrostriatal dopamine system degeneration. Advanced olfactory tests able to measure discrimination are now available for use in rats and could be tested in the rotenone model (Tillerson et al. 2006). More detailed sleep studies need to be performed on rats. In particular, given the risk of developing PD in those with RBD, particular focus should be on REM and muscle activity to assess atonia during REM. Evaluation should be repeated in conjunction with other behavioral assays to determine when sleep disturbances arise relative to motor deficits.

# 5.2 What Are Some Potential Future Uses of Preclinical Rotenone Model?

The identification of preclinical features in the rotenone model provides further validation of model utility. The key question is as follows: how can modeling preclinical features of PD be used to advance our understanding of the disease? It is important to note that although it is now over a decade in use, the rotenone model is still several decades younger than other prominent neurotoxicant modes (6-OHDA, MPTP). Thus, we are still learning much about the model.

In humans, imaging can identify neurological dysfunction prior to diagnosis. Dopaminergic PET and SPECT have high sensitivity and specificity for parkinsonism, with studies estimating that measurable abnormalities may occur 4–7 years before clinical PD (Hilker et al. 2005; Morrish et al. 1998; Postuma et al. 2010; Ravina et al. 2005; Vingerhoets et al. 1994). As these imaging techniques become readily available for rodents with high enough resolution, it should be possible to noninvasively determine temporally, when dopaminergic dysfunction begins in live animals. This would be particularly useful in low-dose, long-term studies, where sampling for biomarkers and nondopaminergic effects can be obtained. Here, an emphasis on what occurs prior to or near the beginning of nigrostriatal dopamine system dysfunction will be important.

Most PD cases likely arise from multiple insults. Thus, when studying etiological factors, exposure to environmentally relevant mixtures at different times in development/life span and exploring the effects of gene-environment interactions are important. Practically, in a laboratory setting, this is expensive and difficult, because even studying a single insult in an animal model is costly and time consuming. Nonetheless, combination and multiple "hit" approaches are being explored in animal models and are becoming more common as a way to address exposures relevant to human health. Given its environmental relevance and now being implicated as a bona fide risk factor, rotenone exposure is particularly relevant to such studies. For example, low-dose rotenone (2.0–2.5 mg/kg/day) reproduces many preclinical features as described above, yet in the absence of overt nigrostriatal toxicity. This regimen is ideal for examining potentiation and synergistic effects from other insults. A recent study where striatal dopamine terminal loss and severe parkinsonian behavioral features were observed in rats expressing disease-causing mutations in  $\alpha$ -synuclein, but not in wild type, shows how the model can be used to assess potentiation (Cannon et al. 2013). It is expected that such forms of the rotenone model will continue to be used in transgenic animals or along with other toxicants of relevance to continue to identify etiological factors.

# 6 Conclusion

The rotenone model replicates the key behavioral, neurochemical, and pathological features of PD. Many models, both neurotoxicant and genetic based, have shown promise in replicating at least some of the key preclinical features of PD. Here, we have presented specific evidence that the rotenone model replicates many key preclinical features of human PD. It has the major advantages of environmental relevance, chronic development, and reproduction of important pathological hall-marks. Future studies are required, but it is expected that findings from the rotenone model will confirm its relevance to model preclinical features that are not yet known to occur in human PD patients.

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Part III

# **Glutamate Receptor Agonists and Antagonists**

# **Concept of Excitotoxicity via Glutamate Receptors**

# Juan C. Piña-Crespo, Sara Sanz-Blasco, and Stuart A. Lipton

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#### Abstract

Since its inception, the concept of glutamate excitotoxicity has provided a foundational framework for understanding the role played by excitatory amino acids in disease states of the brain. At the same time, it has served as a guiding principle in the development and evaluation of new anti-excitotoxic drugs, many of which show promise as neuroprotective therapies in a number of neurological conditions. The discovery that glutamate receptors on the cell surface can engage, through second messengers such as calcium (Ca<sup>2+</sup>), nitric oxide (NO), and inositol

e-mail: pina-crespo@sbmri.org; slipton@sbmri.org

J.C. Piña-Crespo (🖂) • S. Sanz-Blasco • S.A. Lipton

Del E. Webb Center for Neuroscience, Aging, and Stem Cell Research, Sanford-Burnham Medical Research Institute, La Jolla, CA, USA

phospholipids, downstream intracellular signaling cascades involved in cell death helped uncover the complexity of the excitotoxic cascade. The identification of numerous intracellular effectors of excitotoxicity has provided a physiological and pharmacological basis for understanding the cellular and molecular mechanisms behind glutamate-mediated nerve cell injury and its role in neuropsychiatric diseases. More recently, knowledge of the molecular biology of glutamate receptors has allowed, for the first time, the identification of differences in the pattern of expression of glutamate receptors in human populations afflicted by neuropsychiatric diseases. This knowledge will be useful in uncovering genes that may confer individual susceptibility to excitotoxic damage and, as a result, predisposition to the development of certain mental and neurological diseases. In this chapter, the role of glutamate receptor overactivation in excitotoxic cell injury as well as potential neuroprotective therapies for limiting glutamate-mediated neurotoxicity in disease states of the central nervous system will be discussed.

List of Abb	previations
AMPA	2-amino-3-(3-hydroxy-5-methyl-isoxazol-4-yl)propanoic acid)
ATP	Adenosine-5'-triphosphate
BAD	Bcl-2-associated death promoter
BAX	Bcl-2-associated X protein
Bcl-2	B-cell lymphoma 2 protein
$Bcl-X_L$	B-cell lymphoma – extra large protein
Ca <sup>2+</sup>	Calcium
DA	Domoic acid
DCD	Delayed calcium deregulation
DNA	Deoxyribonucleic acid
EAAs	Excitatory amino acids
ER	Endoplasmic reticulum
GAPDH	Glyceraldehyde-3-phosphate dehydrogenase
Glu	L-glutamate
GluRs	Glutamate receptors
KA	Kainic acid
MCU	Mitochondrial calcium uniporter
mNCX	Mitochondrial Na <sup>+</sup> /Ca <sup>2+</sup> transporter
mPTP	Mitochondrial permeability transition pore
NCX	Na <sup>+</sup> /Ca <sup>2+</sup> exchanger
NMDA	N-methyl-D-aspartic acid
NO	Nitric oxide
PMCA	Plasma membrane Ca <sup>2+</sup> ATPase
RNS	Reactive nitrogen species
ROS	Reactive oxygen species
SIAH1	Seven in absentia homolog E3 ubiquitin-protein ligase
VDAC	Voltage-dependent anion-selective channel
$\Delta \psi m$	Inner mitochondrial membrane potential

### 1 Introduction

L-Glutamate is a major excitatory amino acid neurotransmitter in the brain (Nicholls et al. 2012; Watkins and Evans 1981; Watkins 2000). Under physiological conditions, glutamate plays fundamental roles in nerve cell function and brain plasticity (Bliss and Collingridge 1993; Cooke and Bliss 2006) through activation of glutamate receptors (GluRs) (Hollmann and Heinemann 1994; Traynelis et al. 2010). In contrast, excessive activation of GluRs during neurological conditions can lead to loss of nerve cell function and cell death through a process known as excitotoxicity (Olney 1978; Olney 2003). Glutamate-mediated excitotoxicity is initiated by buildup of excess glutamate in the extracellular space that leads to overactivation of GluRs, disruption in cell membrane permeability, and downstream activation of signaling cascades involved in loss of nerve cell function, cell damage, and death (Farooqui et al. 2008; Gillessen et al. 2002; Lau and Tymianski 2010). Significant experimental and clinical evidence implicates excitotoxicity in the loss of neuronal cell function and cell death associated with acute neurological conditions such as traumatic brain injury (Arundine and Tymianski 2004; Yi and Hazell 2006), stroke (Besancon et al. 2008; Brouns and De Deyn 2009; Doyle et al. 2008), hypoxiaischemia (Jensen 2005), amnesic shellfish poisoning (Jeffery et al. 2004; Lefebvre and Robertson 2010; Perl et al. 1990; Teitelbaum et al. 1990), and epileptic seizures (Meldrum 1993), as well as during chronic neurodegenerative diseases (Lipton and Rosenberg 1994; Mattson 2003; Mehta et al. 2013) such as Alzheimer's disease (Hu et al. 2012; Mucke and Selkoe 2012), Parkinson's disease (Beal 1998; Surmeier et al. 2011), Huntington's disease (Raymond et al. 2011), AIDS encephalopathy and dementia complex (more recently termed HIV-associated neurocognitive disorders) (Kaul et al. 2001; Mattson et al. 2005), multiple sclerosis (Trapp and Stys 2009), and neuropathic pain syndromes (Latremoliere and Woolf 2009).

Neuronal function, as well as susceptibility to excitotoxic cell damage, depend on the expression of specific repertoires of receptors and ion channels. Since abnormal patterns of expression of genes encoding glutamate receptors have been described in a number of brain disorders, current research efforts may unveil how dysregulated expression of glutamate receptors affects neurotransmission and nerve cell susceptibility to excitotoxic cell death.

### 2 Historical Background

Although interest into the role of glutamate in the nervous system dates back to the early twentieth century (Weil-Malherbe 1950), it was not until the 1950s that evidence for a direct excitatory action of glutamate on the brain was first described (Hayashi 1952; Okamoto 1951) and insights into its mechanism of action were first inferred (Hayashi 1954). These findings were soon followed by reports of glutamate-induced neurotoxicity after observation of acute degenerative lesions in the inner retina of newborn mice (Lucas and Newhouse 1957) and depolarization of neurons in the spinal cord (Curtis et al. 1959).

During the 1960s, research into the structure-function relationship of a large number of neutral and acidic amino acids led to the concept an "amino-acid-receptor complex" present in the extracellular surface of neurons (Curtis et al. 1960). In an effort to elucidate the chemical basis of the activity and specificity of glutamate and aspartate on the "amino acid receptor," Watkins and colleagues synthesized *N*-methyl-D-aspartic acid (NMDA) (Watkins 1962), a non-metabolizable agonist with remarkable selectivity for the "amino-acid-receptor complex." Sensitivity to NMDA became a key criterion for the initial pharmacological classification of glutamate receptors into NMDA and non-NMDA receptors (Lodge 2009).

Further evidence supporting a neurotoxic role for glutamate appeared in 1969, when Olney and collaborators described neuropathological lesions in the brain of rodents and primates after systemic administration of glutamate and structurally related analogues (Olney 1969; Olney and Sharpe 1969). During the ensuing decade, Olney and colleagues proposed the concept of excitotoxicity to describe the acute neurotoxic effects of glutamate in the brain (Olney and Ho 1970; Olney 1978) and provided further experimental support for its potential role in brain pathology particularly in the newborn brain.

During the 1980s, preclinical studies reported evidence of significantly high levels of extracellular concentrations of brain glutamate in experimental models of hypoxia-ischemia in vivo (Benveniste et al. 1984; Hagberg et al. 1985), a finding later confirmed in humans (Bullock et al. 1995). At the same time, preclinical studies confirmed the neuroprotective action of glutamate receptor antagonists against hypoxic-ischemic neuronal cell death in vitro (Rothman 1984) and in vivo (Simon et al. 1984). However, successful translation of these early preclinical findings into clinically effective therapies has been an elusive target (O'Collins et al. 2006; Sutherland et al. 2012; Tymianski 2010), mainly because of lack of therapeutic strategies that can selectively target pathologically active GluRs without affecting normal glutamatergic neurotransmission (Lipton 2007).

A need for understanding the intracellular signaling cascades activated by neurotoxic glutamate led to the identification of calcium (Ca<sup>2+</sup>) (Choi 1985; Choi 1987), inositol phospholipids (Sladeczek et al. 1985; Sugiyama et al. 1987), and nitric oxide (NO) (Garthwaite et al. 1988) as key intracellular effectors of glutamate signaling. This set of key insights provided the physiological and pharmacological frameworks for a greater understanding of the cellular and molecular mechanisms responsible for glutamate-mediated nerve cell injury (Choi 1988; Rothman and Olney 1986) and its potential role in neurological diseases (Greenamyre 1986). In 1987, a fatal incident due to consumption of shellfish contaminated with domoic acid (DA), a naturally occurring glutamate analogue, provided direct evidence for glutamate-mediated neurotoxicity in humans (Perl et al. 1990; Teitelbaum et al. 1990).

During the 1990s, glutamate research experienced one of its most important and significant periods with the cloning and functional expression of 26 different glutamate receptor genes (Table 1) (Alexander et al. 2011; Boulter et al. 1990; Hollmann et al. 1989; Hollmann and Heinemann 1994). This unveiled the molecular basis for the functional and pharmacological diversity of glutamate receptors in the central nervous system (Gereau and Swanson 2008; Traynelis et al. 2010).

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	Receptor	Pharmacology	Gene		Previous nomenc	lature
Ionotropic receptors			Human	Mouse/rat	Rat	Mouse
1	GluA1	AMPA	GRIA1	Gria1	GluR1, GluRA	α1
2	GluA2	AMPA	GRIA2	Gria2	GluR2, GluRB	α2
3	GluA3	AMPA	GRIA3	Gria3	GluR3, GluRC	α3
4	GluA4	AMPA	GRIA4	Gria4	GluR4, GluRD	α4
5	GluD1	Glutamate	GRID1	Grid1	Delta1	δ1
6	GluD2	Glutamate	GRID2	Grid2	Delta2	δ2
7	GluK1	Kainate	GRIK1	Grik1	GluR5, GluK5	β1
8	GluK2	Kainate	GRIK2	Grik2	GkuR6, GluK6	β2
9	GluK3	Kainate	GRIK3	Grik3	GluR7, GluK7	β3
10	GluK4	Kainate	GRIK4	Grik4	KA1, GluK1	γ1
11	GluK5	Kainate	GRIK5	Grik5	KA2, GluK2	γ2
12	GluN1	NMDA	GRIN1	Grin1	NR1	ζ1
13	GluN2A	NMDA	GRIN2A	Grin2a	NR2A	ε1
14	GluN2B	NMDA	GRIN2B	Grin2b	NR2B	ε2
15	GluN2C	NMDA	GRIN2C	Grin2c	NR2C	ε3
16	GluN2D	NMDA	GRIN2D	Grin2d	NR2D	ε4
17	GluN3A	NMDA	GRIN3A	Grin3a	NR3A	χ1
18	GluN3B	NMDA	GRIN3B	Grin3b	NR3B	χ2
Met	abotropic rec	eptors				
19	mGlu <sub>1</sub>	Glutamate	GRM1	Grm1	mGluR1	mGluR
20	mGlu <sub>2</sub>	Glutamate	GRM2	Grm2	mGluR2	mGluR
21	mGlu <sub>3</sub>	Glutamate	GRM3	Grm3	mGluR3	mGluR
22	$mGlu_4$	Glutamate	GRM4	Grm4	mGluR4	mGluR
23	mGlu <sub>5</sub>	Glutamate	GRM5	Grm5	mGluR5	mGluR
24	mGlu <sub>6</sub>	Glutamate	GRM6	Grm6	mGluR6	mGluR
25	mGlu <sub>7</sub>	Glutamate	GRM7	Grm7	mGluR7	mGluR
26	mGlu <sub>8</sub>	Glutamate	GRM8	Grm8	mGluR8	mGluR

**Table 1** IUPHAR\* classification of cloned glutamate receptors (GluRs)

\*IUPHAR Committee on Receptor Nomenclature and Drug Classification (http://www.iuphar-db. org/index.jsp)

The molecular biology of glutamate receptors laid down the foundations for detailed cellular, molecular, and genetic studies into the role of glutamate dysfunction in diseases of the brain (Lipton and Rosenberg 1994; Nakanishi 1992).

### **3 Glutamate Receptors**

Glutamate receptor-mediated signaling occurs through activation of two different types of receptors: ion channel-coupled (ionotropic) and G-protein-coupled (metabotropic) receptors. Ionotropic glutamate receptors (iGluRs) are coupled to ion channels and are responsible for fast changes in membrane permeability

(Traynelis et al. 2010; VanDongen 2009); metabotropic glutamate receptors (mGluRs) are coupled to intracellular effector systems via G-proteins (Nicoletti et al. 2011; Niswender and Conn 2010).

### 3.1 Ionotropic Glutamate Receptors

iGluRs are essential mediators of fast synaptic transmission due to their ability to generate a response within milliseconds of glutamate binding. Based on their agonist selectivity, iGluRs have been classified into three major subclasses: AMPA ( $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazole propionic acid), kainate, and NMDA receptors (Table 1). iGluRs are cation-selective glutamate-gated ion channels composed of four glycosylated transmembrane subunits assembled around a central aqueous pore that spans the cell membrane. Each receptor contains three major domains: an extracellular glutamate-binding domain related to amino acid-binding bacterial periplasmic proteins, a transmembrane pore-lining domain related to the pore-forming region or "P-loop" of K<sup>+</sup> channels, and an intracellular regulatory domain (Gereau and Swanson 2008; Kumar and Mayer 2012; Traynelis et al. 2010).

### 3.1.1 AMPA Receptors

Upon activation by L-glutamate or AMPA, AMPARs mediate responses that are characterized by rapid onset, offset, and desensitization kinetics, but when they are activated by the neurotoxin kainate or domoic acid (DA), they produce large non-desensitizing responses. Four AMPAR subunits have been cloned and termed: GluA1, GluA2, GluA3, and GluA4 (Alexander et al. 2011; Boulter et al. 1990; Keinanen et al. 1990); they can form homomeric receptors as well as heteromeric receptors when expressed with other subunits. Functionally, they show rapid kinetics and low  $Ca^{2+}$  permeability in the presence of the GluA2 subunit (although the RNA-unedited form of GluA2 is permeable to  $Ca^{2+}$ ) (Gereau and Swanson 2008; Traynelis et al. 2010).

### 3.1.2 Kainate Receptors

So far, five different sequence-related genes encoding kainate receptors (KARs) have been identified and termed: GluK1, GluK2, GluK3, GluK4, and GluK5. KARs bind kainate with intermediate to high affinity with GluK1 and GluK2 being the only kainate receptors capable of forming functional receptors upon in vitro expression. The others kainate receptor subunits can form functional receptors only when co-expressed in combination with GluK1 or GluK2 receptors. In contrast with AMPARs, KARs strongly desensitize upon activation by the neurotoxin kainate.

### 3.1.3 NMDA Receptors

NMDA receptors are characterized by high  $Ca^{2+}$  permeability, voltage-dependent block by  $Mg^{2+}$ , high single-channel conductance, slow gating kinetics, and activation by the concerted action of glutamate and glycine (VanDongen 2009).

During glutamatergic synaptic transmission, NMDA receptors mediate the slow component of synaptic currents. NMDA receptors result from the assembly of three different subfamilies of subunits: NR1, NR2, and NR3 (Table 1).

# 3.2 Metabotropic Glutamate Receptors

Metabotropic glutamate receptors (mGluRs) play an important role in the transduction of glutamate-mediated signaling via coupling through G-proteins to intracellular second messenger cascades. They have been classified into groups 1, 2, and 3. Group 1 (mGluR1 and mGluR5) receptors are coupled to phospholipase C, mediating increases in intracellular calcium. Group 2 (mGluR2 and mGluR3) and group 3 (mGluR4, mGluR6, mGluR7, mGluR8) are negatively coupled to adenylate cyclase, producing a decrease in forskolin-induced cAMP (Nicoletti et al. 2011; Niswender and Conn 2010).

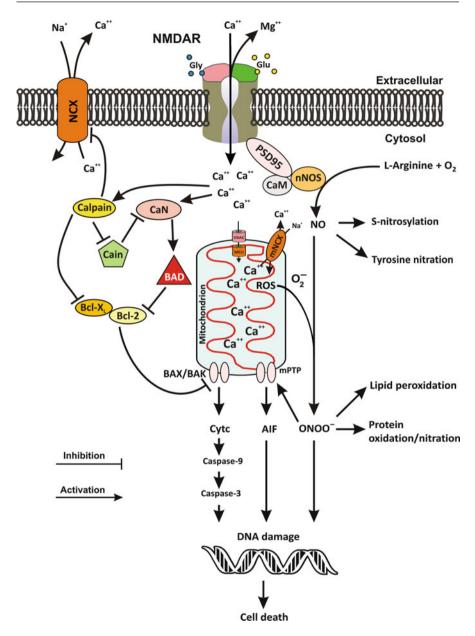
# 4 Mechanisms of Excitotoxicity

Glutamate neurotoxicity results from a buildup of either endogenous excitatory amino acids (EAAs), exogenous glutamate-like substances in the brain, or excessive glutamate receptor activity (often NMDA receptors) in the presence of normal agonists due to loss of normal  $Mg^{2+}$  block. This form of neurotoxicity is often due to malfunctioning of cellular processes responsible for regulating endogenous EAA levels in the brain or to exposure to toxic levels of exogenous EAA analogues known as excitotoxins. These events produce intense activation of glutamate receptors on the cell surface that generates abnormally high levels of intracellular  $Ca^{2+}$ , which activates catalytic enzymes, generates highly toxic free radicals, and disrupts cell energy production. Eventual depletion of ATP stores leads to the collapse of transmembrane electrochemical gradients, loss of nerve cell excitability, neurotransmission failure, and cell death by either necrosis or apoptosis (Choi 1988; Gillessen et al. 2002; Rothman and Olney 1986).

# 5 Intracellular Effectors of Glutamate Neurotoxicity

## 5.1 Calcium

 $Ca^{2+}$  ions are powerful activators of nerve cell excitability and metabolism. Intracellular  $Ca^{2+}$  concentrations are kept relatively low by high-energy-requiring processes of active  $Ca^{2+}$  extrusion and  $Ca^{2+}$  sequestration in intracellular stores. Active  $Ca^{2+}$  efflux is mostly carried out by the Na<sup>+</sup>/Ca<sup>2+</sup> exchanger (NCX) and plasma membrane  $Ca^{2+}$  ATPase (PMCA), while storage of intracellular  $Ca^{2+}$  localizes mainly to the endoplasmic reticulum (ER) and mitochondria. Extracellular  $Ca^{2+}$ concentrations (1–2 mM) are over one order of magnitude larger than inside nerve



**Fig. 1** Intracellular effectors of excitotoxicity. The events leading to excitotoxic cell death include overactivation of  $Ca^{2+}$ -permeable glutamate receptors (GluRs), excessive  $Ca^{2+}$  influx, and  $Ca^{2+}$ -dependent activation of intracellular effector enzymes such neuronal nitric oxide synthase (nNOS), calcineurin (CaN), and calpain together with  $Ca^{2+}$  uptake into mitochondria. Subsequent mitochondrial dysfunction due to excessive  $Ca^{2+}$  accumulation, and generation and release of toxic amounts of free radicals such as nitric oxide (NO) and reactive oxygen species (ROS) contribute to the production of peroxynitrite (ONOO<sup>-</sup>) that together with depletion of

cells (50–200 nM). During overactivation of Ca<sup>2+</sup>-permeable GluRs (Brorson et al. 1994; Choi 1985; Lu et al. 1996), this large electrochemical gradient drives the flow of extracellular Ca<sup>2+</sup> into the cytoplasm where it generates an over tenfold increase in intracellular Ca<sup>2+</sup>. In vitro, even a brief, few seconds long, exposure to neurotoxic levels of glutamate can trigger loss of intracellular Ca<sup>2+</sup> homeostasis and cell death. Glutamate-induced intracellular loading of Ca<sup>2+</sup> is characterized by an immediate but transient GluR-mediated Ca<sup>2+</sup> influx that is followed by a transient return to baseline Ca<sup>2+</sup> levels that can last hours until a secondary wave of sustained intracellular Ca<sup>2+</sup> develops that is irreversibly followed by cell death (Randall and Thayer 1992). This dynamic demonstrates how GluR-mediated calcium influx activates specific effector pathways involved in cell death. Intense excitotoxic injury can result in necrosis (Syntichaki and Tavernarakis 2003), while less fulminant or chronic insults can cause synaptic damage followed by apoptosis (Ankarcrona et al. 1995; Bonfoco et al. 1995).

Although there is general consensus regarding the important role of intracellular  $Ca^{2+}$  in excitotoxic nerve cell death (Choi 1985; Choi 1988), many of the intracellular signaling events underlying glutamate-induced cell death are less well characterized. Reactive oxygen species (ROS), reactive nitrogen species (RNS), caspases, and lipases are among the most important  $Ca^{2+}$ -dependent effectors thought to be involved in glutamate-induced neurotoxicity. These processes are detailed below and summarized in Fig. 1.

# 5.2 Reactive Oxygen Species (ROS)

GluR-mediated excitotoxicity generates reactive oxygen species (ROS) such as superoxide  $(O_2^-)$ , hydrogen peroxide  $(H_2O_2)$  and hydroxyl (OH<sup>-</sup>) radicals (Dykens et al. 1987; Dykens 1994; Lafon-Cazal et al. 1993). ROS cause cell damage through lipid peroxidation and oxidative damage of proteins and DNA (Chinopoulos and Adam-Vizi 2006; Rigoulet et al. 2011; Van Houten et al. 2006). Numerous neuroprotective strategies, including trapping, scavenging, and inhibition of ROS synthesis, have been shown to be effective under in vitro conditions (Dykens et al. 1987; Majewska and Bell 1990; Puttfarcken et al. 1992); however, evidence of their

**Fig. 1** (continued) mitochondrial ATP levels promotes opening of the BAX/BAK channel, mitochondrial permeability transition pore (mPTP), or breakdown of the mitochondrial membrane to release cytochrome c (Cyt c) and the apoptosis-inducing factor (AIF). While Cyt c mediates caspase-dependent apoptosis, AIF translocates into the nucleus where it produces DNA fragmentation, chromatin condensation, and apoptosis through a caspase-independent mechanism. Simultaneously,  $Ca^{2+}$ -dependent activation of calcineurin (CaN) and calpain promotes calpain-dependent inhibition of the anti-apoptotic Bcl-X<sub>L</sub>/Bcl-2 complex and Cain, an inhibitor of calcineurin-mediated cell death. Once calcineurin (CaN) is freed from Cain-dependent inhibition, it can activate the Bcl-2-associated death promoter (BAD), which produces further inhibition of Bcl-X<sub>L</sub>/Bcl-2 complex. This allows the formation of a BAX/BAK mitochondrial channel permeable to pro-apoptotic molecules such as Cyt c

neuroprotective activity in vivo is still lacking. Since mitochondria are both a source of ROS (Dugan et al. 1995; Reynolds and Hastings 1995) and a target for ROS-induced damage, protection and rescue of mitochondrial function is key to reducing glutamate-mediated neurotoxicity (Cho et al. 2010; Perez-Pinzon et al. 2012).

### 5.3 Reactive Nitrogen Species (RNS)

Nitric oxide (NO) can react with superoxide radicals  $(O_2^{-})$  to generate peroxynitrite (ONOO<sup>-</sup>), a highly toxic and strongly oxidizing and nitrating agent that causes lipid peroxidation, protein oxidation, and DNA damage (Brown 2010). Inhibition of NO synthesis or genetic deletion of its synthesizing enzyme, nitric oxide synthetase (nNOS), can protect neurons from glutamate neurotoxicity (Dawson et al. 1991; Dawson et al. 1996; Lipton et al. 1993). Because of its close physical association to intracellular scaffoldings such as postsynaptic densities (PSDs), which anchor GluRs to postsynaptic sites (Sattler et al. 1999), nNOS is one of the first Ca<sup>2+</sup>-dependent intracellular effectors activated by Ca<sup>2+</sup> influx through GluRs, namely, NMDARs. Through this pathway, neurotoxic glutamate levels generate large amounts of RNS, especially NO, which causes misfolding and malfunction of intracellular signaling proteins through abnormal S-nitrosylation and tyrosine nitration (Martinez-Ruiz et al. 2011; Nakamura and Lipton 2011). Among the most important deleterious effects of NO accumulation is inhibition of energy-generating intracellular systems that leads to the collapse of intracellular ATP levels and cell death through either necrosis or apoptosis. During this process, NO causes further release of glutamate through inhibition of mitochondrial function, reverse glutamate uptake, and glutamate exocytosis (Brown 2010). In addition, neuronal cell death can be triggered by NO-dependent nitrosylation of the glycolytic enzyme glyceraldehyde-3-phosphate dehydrogenase (GAPDH), which promotes its association with the E3 ubiquitin-protein ligase SIAH1 and translocation into the nucleus (Hara et al. 2005). There is currently considerable need for effective cell-permeable inhibitors to protect nerve cells from the abnormally high levels of protein nitrosylation observed during pathological overactivation of brain GluRs (Nakamura and Lipton 2010).

### 5.4 Caspases

Caspases are intracellular aspartate-specific proteases that play a central role in apoptotic nerve cell death (Troy et al. 2011). They are strongly associated with glutamate-induced excitotoxic apoptosis (Tenneti et al. 1998; Tenneti and Lipton 2000), where their activation occurs via the mitochondrial or intrinsic apoptotic pathway (Budd et al. 2000). Overactive glutamate receptors produce excessive calcium influx and ROS, which causes mitochondrial calcium overload and energy depletion that triggers mitochondrial permeabilization and release of

cytochrome c (Cyt *c*) into the cytoplasm. Cyt *c* relieves calcium-induced inhibition of inositol trisphosphate receptors ( $IP_3R$ ) and initiates the formation of the apoptosome which activates the apoptotic caspase cascade (Kroemer et al. 2007). The interplay between activated caspases and calpains is thought to underlie the neurotoxic effect of glutamate through excitotoxic apoptosis (Hara and Snyder 2007).

#### 5.5 Calpains

Calpains are a group of intracellular Ca<sup>2+</sup>-dependent cysteine proteases activated during excitotoxic apoptosis (Ankarcrona et al. 1995; D'Orsi et al. 2012; Vosler et al. 2008). Calcium overload due to GluR overactivation causes calpain activation and subsequently cell death through cleavage and inactivation of signaling and structural intracellular proteins (Brorson et al. 1995; Liu et al. 2008; Siman et al. 1989). Although the exact sequence of events is not clearly understood, the interplay between caspases and calpains (Wu et al. 2007) involves Ca<sup>2+</sup>-dependent activation of calpain which causes cleavage and internalization of the membranebound Na<sup>+</sup>/Ca<sup>2+</sup> exchanger 3 (NCX3) and leads to an increase in intracellular calcium and cell death in a form of active necrosis (Bano et al. 2005). Activated calpains also target the protein Cain/Cabin-1, an intracellular inhibitor of calcineurin-mediated cell death (Kim et al. 2002; Lai et al. 1998). Calpain-mediated Cain/Cabin-1 inactivation removes a key restrictive control over calcineurin dephosphorylation of the pro-apoptotic protein Bcl-2-associated death promoter (BAD). Dephosphorylated BAD binds and inactivates the antiapoptotic proteins Bcl-2 and Bcl-X<sub>L</sub> (Wang et al. 1999). In addition, activated calpains can directly cleave and inactivate Bcl-2 and Bcl-X<sub>L</sub> (Vandenabeele et al. 2005), which are responsible for preventing activation of the pro-apoptotic mitochondrial membrane permeabilization complex formed by Bcl-2-associated X protein (BAX)/Bcl-2antagonistic/killer (BAK) (Yang et al. 1997). Activation of the BAX/BAK complex forms a pore on the mitochondrial outer membrane that is responsible for the release of cytochrome c that initiates caspase-mediated apoptosis (Liu et al. 1996; Ow et al. 2008). An alternative pathway activated by calpains during excitotoxicity involves activation and release of the apoptosis-inducing factor (AIF) (Cao et al. 2007; Landshamer et al. 2008; Polster et al. 2005; Yu et al. 2002), a mitochondrial flavoprotein involved in caspase-independent apoptosis (Susin et al. 1996). The importance of calpain inhibitors as a potential neuroprotective agents is highlighted by their significant neuroprotective effect against excitotoxic brain damage induced by cerebral ischemia and traumatic brain injury (Koumura et al. 2008; Kupina et al. 2001; Lubisch et al. 2003; Nimmrich et al. 2010; Wang et al. 1996).

### 5.6 Lipases

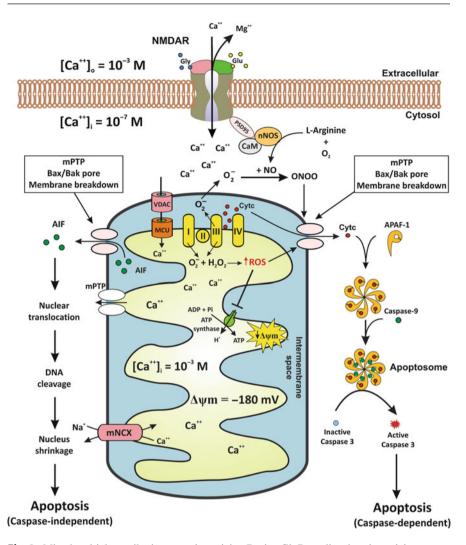
NMDAR-mediated glutamate neurotoxicity activates  $Ca^{2+}$ -dependent phospholipase A2 (PLA2), which triggers release of arachidonic acid (AA) from membrane phospholipids (Dumuis et al. 1988; Lazarewicz et al. 1988). Extracellular release of arachidonic acid can inhibit transporter-mediated glutamate (Volterra et al. 1992), while intracellular metabolism of AA by lipoxygenases (LOX), cytochrome P450 (CYP), and cyclooxygenases (COX) generates bioactive eicosanoids that trigger neuroinflammatory cell damage.

#### 6 Role of Mitochondria in Glutamate Excitotoxicity

Calcium accumulation in mitochondria is a key event in glutamate excitotoxicity (Duchen 2012; Orrenius et al. 2003). It has been shown in many neurodegenerative diseases that overactivation of calcium-permeable GluRs causes an increase of intracellular calcium that induces overloading of calcium in the mitochondria, followed by an increase in free radicals and a decrease in ATP levels (Thayer and Wang 1995). In fact, the amount of calcium taken up by the mitochondria is decisive in cell death by glutamate excitotoxicity. Once the mitochondrial capacity for Ca<sup>2+</sup> sequestration is exceeded due to depletion of mitochondrial energy resources, mitochondrial failure occurs (Rizzuto et al. 2012) (Fig. 2).

Physiologically, Ca<sup>2+</sup> transport to the mitochondria through a complex formed between the voltage-dependent anion-selective channel (VDAC) and the mitochondrial calcium uniporter (MCU) contributes to the maintenance of low levels of Ca<sup>2+</sup> in the cytosol (Rizzuto et al. 2012). The main route of  $Ca^{2+}$  entry into the mitochondria is through the  $Ca^{2+}$  uniporter (MCU), which is activated by  $Ca^{2+}$  in an allosteric manner (Kirichok et al. 2004). During pathological increases in cytosolic Ca<sup>2+</sup> concentration, mitochondria accumulate Ca<sup>2+</sup> ions and sometimes Ca<sup>2+</sup>phosphate complexes form as reversible calcium deposits (David et al. 2003; Pivovarova et al. 1999). In excitable cells, the primary method of calcium extrusion out of the mitochondria is the  $Na^+/Ca^{2+}$  transporter (mNCX), responsible of the low basal [Ca<sup>2+</sup>]<sub>mit</sub>. Because of its low carrier capacity, the Na<sup>+</sup>/Ca<sup>2+</sup> transporter (mNCX) can be easily overtaken by  $Ca^{2+}$  influx through the  $Ca^{2+}$  uniporter (MCU), generating a net accumulation of  $Ca^{2+}$  in mitochondria. An alternative pathway for  $Ca^{2+}$  extrusion is the H<sup>+</sup>/Ca<sup>2+</sup> exchanger, but it is slow and saturates at a lower  $[Ca^{2+}]_{Cit}$  than the Na<sup>+</sup>/Ca<sup>2+</sup> transporter; so its contribution to fast calcium exchange is limited (Bernardi 1999; Gunter et al. 1991).

Still another route for  $Ca^{2+}$  extrusion is via the transitory opening of the mitochondrial permeability transition pore (mPTP) complex, which controls excessive influx of toxic levels of  $Ca^{2+}$  (Bernardi and von Stockum 2012; Ichas et al. 1997; Zoratti and Szabo 1995). It is known that factors such as increased oxidative stress (ROS), mitochondrial  $Ca^{2+}$  overload, and metabolic stress promote mPTP opening (Duchen 2000a; Duchen 2000b). mPTP opening determines whether a cell dies or survives. Although its opening mechanism is not very well understood, it is known that mitochondrial  $Ca^{2+}$  overload is enough to promote its opening. It has been proposed that partial mitochondrial depolarization can prevent  $Ca^{2+}$  overload in mitochondria (Budd and Nicholls 1996; Sanz-Blasco et al. 2008). Simultaneous measurements of  $[Ca^{2+}]$  and inner mitochondrial



**Fig. 2** Mitochondrial contribution to excitotoxicity. During GluR-mediated excitotoxicity, excess  $Ca^{2+}$  is transported into mitochondria by a complex formed between the voltage-sensitive anion channel (VDAC) and the mitochondrial  $Ca^{2+}$  uniporter (MCU). Although  $Ca^{2+}$  concentrations are higher inside the mitochondria compared with the cytosol,  $Ca^{2+}$  is normally transported inside mitochondrial membrane potential ( $\Delta \psi m = -180 \text{ mV}$ ). As excess  $Ca^{2+}$  accumulates, it produces a persistent depolarization of the inner mitochondrial membrane potential that triggers the synthesis of ATP via the ATP synthase and increases the activity of the electron transport chain with the subsequent production of reactive oxygen species (ROS). Persistent accumulation of ROS in mitochondria may inhibit ATP synthesis and the activity of the respiratory chain, leading to energy failure and loss of mitochondrial homeostasis. As the integrity of the mitochondria deteriorates, key internal components of the mitochondria such as cytochrome c (Cyt c) and apoptosis-inducing factor (AIF) are released into the cytosol where they act as powerful triggers of apoptosis. Cyt c interacts with apoptotic protease-activating factor 1 (APAF-1) and

membrane potential ( $\Delta \psi m$ ) in hippocampal neurons during exposure to toxic glutamate levels show that [Ca<sup>2+</sup>]c increases transiently followed by a variable degree of recovery. This is then followed by a failure of the cell to maintain a low cytoplasmic free Ca<sup>2+</sup> concentration, called delayed calcium deregulation (DCD). During this process,  $\Delta \psi m$  changes only slightly during the initial phase, showing a progressive and complete depolarization later on (Abramov and Duchen 2008) that causes ATP depletion and cell death. GluR-dependent transient mitochondrial calcium loading can also initiate oxidative damage and/or inhibit mitochondrial respiration (Ward et al. 2000) observed as a reduction in oxygen consumption due to inhibition of mitochondrial complexes I, II/III, and IV (Rego et al. 2000).

## 7 Clinical Relevance of GluR-Mediated Excitotoxicity

Although the original concept behind the mechanism of excitotoxicity originated in strong preclinical evidence suggesting that acute and chronic neurotoxicity can use the excitatory amino acid glutamate to produce nerve cell injury and cell death (Olney 1969; Olney 1978), it was not until 20 years later that a direct link between glutamate excitotoxicity and clinical disease in humans could be identified (Perl et al. 1990; Teitelbaum et al. 1990). Although equally strong preclinical (Benveniste et al. 1984; Cook et al. 2012a; Cook et al. 2012b; Hagberg et al. 1985) and clinical (Bullock et al. 1995) evidence supports a significant role for GluR-mediated excitotoxicity in hypoxia-ischemia and stroke, the initial pace of discovery of anti-excitotoxic drugs based in GluR antagonists has been slower than expected (O'Collins et al. 2006; Savitz and Fisher 2007). One of the reasons for the slow progress of discovery of antiexcitotoxic therapies based in GluR antagonists has been the fact that most inhibitors of GluRs cannot discriminate between GluRs engaged in physiologically relevant glutamatergic neurotransmission and pathologically active GluRs associated with disease states (Lipton 2004; Lipton 2007). The search for clinically tolerated GluR antagonists that block excitotoxicity by preferentially targeting pathologically active GluRs led to the identification of memantine, an uncompetitive NMDA receptor antagonist (Chen et al. 1992; Chen and Lipton 2006) that preferentially blocks pathologically active GluRs and protects neurons against excitotoxic injury (Fig. 3) (Lipton 2007). Memantine's neuroprotective activity has made it a drug of choice in the treatment of chronic neurological diseases such as Parkinson's (Olney et al. 1987; Rabey et al. 1992), dementia with Lewy bodies (Sabbagh et al. 2005), and Alzheimer's disease, for which it is approved by the FDA for clinical use (Fleischhacker et al. 1986; McShane et al. 2006; Reisberg et al. 2003).

**Fig. 2** (continued) caspase 9 to form the apoptosome, which activates caspase 3, a key powerful mediator of apoptotic cell death. Loss of mitochondrial integrity also allows the release of the apoptosis-inducing factor (AIF) into the cytosol where it enters the nucleus and causes DNA fragmentation, chromatin condensation, and nucleus shrinkage through a caspase-independent mechanism of apoptosis

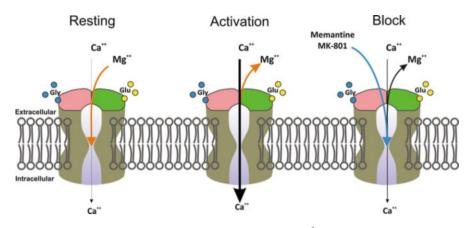


Fig. 3 Preferential block of pathologically active GluRs.  $Ca^{2+}$ -permeable NMDARs are an important mediator of GluR excitotoxicity. Under resting conditions, a steep electrochemical gradient favors Ca<sup>2+</sup> influx through NMDARs while a voltage-dependent block by extracellular Mg<sup>2+</sup> interferes with Ca<sup>2+</sup> influx. During NMDAR activation, transient depolarization is required to force  $Mg^{2+}$  out of the channel and allow  $Ca^{2+}$  influx into the cell. Once the cell returns to its normal resting potential, Mg<sup>2+</sup> block of Ca<sup>2+</sup> influx is restored while excess Ca<sup>2+</sup> is pumped out of the cell. Selective NMDAR antagonists, such as memantine, act as activity-dependent openchannel blockers that, similar to Mg<sup>2+</sup>, can gain access to the inside the ion channel pore and block the influx of Ca<sup>2+</sup> through NMDARs during excessive pathological activation. Because its blocking effect requires prior activation of the receptor by glutamate, which exposes binding sites that are otherwise hidden during the resting state, the efficacy of the drug increases as the activity of the receptor increases and vice versa. In addition to this property, preferential block of pathologically active GluRs requires drugs to be highly selective for their specific target, to avoid interference with other neurotransmitter systems and unwanted side effects, and to have low affinity for its binding site (in the case of memantine, produced by a relatively rapid off-rate from the channel) in order to minimize inhibition of the receptor during normal physiological function. In contrast, MK-801 has a very slow off-rate from the channel, blocking physiological activity and producing multiple side effects

Recently, ketamine, an NMDAR open-channel blocker and dissociative anesthetic with anti-excitotoxic properties (Choi et al. 1989; Choi 1990), has been shown to possess strong antidepressant activity in both preclinical and clinical studies of depression (Skolnick et al. 2009).

An alternative anti-excitotoxic strategy that has produced positive clinical results makes use of drugs that boost the function of glutamate transport systems to reduce extracellular levels of glutamate (Fumagalli et al. 2008; Rothstein 1996). An example of such drugs is riluzole (Fumagalli et al. 2008), which is currently approved by the FDA for the treatment of amyotrophic lateral sclerosis (Bensimon et al. 1994; Lacomblez et al. 1996; Miller et al. 2012).

A growing body of evidence suggests GluR-mediated neurotoxicity facilitates growth of gliomas, one of the most aggressive and malignant forms of brain cancers (Takano et al. 2001; Watkins and Sontheimer 2012; Ye and Sontheimer 1999). Glioma cells can cause excitotoxic cell death in the surrounding brain tissue by releasing toxic amounts of glutamate through the cystine-glutamate antiporter

(SXC, SLC7A11) (Buckingham et al. 2011; Watkins and Sontheimer 2012). Clinical trials are currently under way to test the neuroprotective efficacy of the NMDAR antagonist memantine and the SXC inhibitor sulfasalazine in patients suffering from glioma (*ClinicalTrials.gov*: NCT01260467; NCT01430351, NCT01577966).

#### 8 Conclusion

Although the full implications of excitotoxicity are not yet fully understood, a large body of evidence points towards a role for excitotoxic processes in brain disease. Either as a cause or consequence of other pathological process, the role of excitotoxicity in brain disease has become increasingly undeniable and as a result, difficult to ignore when trying to understand a series of neurodegenerative and neuropsychiatric disorders.

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# Role of Ionotropic Glutamate Receptors in Neurodegenerative and Other Disorders

# Lotten Ragnarsson, Peter R. Dodd, and Matthew R. Hynd

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L. Ragnarsson (🖂)

P.R. Dodd • M.R. Hynd

School of Chemistry and Molecular Biosciences, The University of Queensland St. Lucia Campus, Brisbane, Australia

e-mail: p.dodd@uq.edu.au; m.hynd1@uq.edu.au

Institute for Molecular Bioscience, The University of Queensland, Brisbane, QLD, Australia e-mail: l.ragnarsson@uq.edu.au

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#### Abstract

Disorders of the central nervous system (CNS) are amongst the most complex disease states and lead to some of the most devastating conditions found in medicine today. It is well established that both genetic and epigenetic factors influence the pathogenesis of neurodegenerative disorders. More recently a growing body of evidence suggests that changes in neurotransmitter systems may underlie the pathogenic mechanisms of neurodegenerative disorders, such as Alzheimer's disease and Parkinson's disease. Excitatory amino acids (EAA) are known to play key roles in neurotransmission, neuromodulation, and neurotoxicity. EAA receptors are classified into two major subdivisions, ionotropic glutamate receptors (iGluRs) and metabotropic glutamate receptors (mGluRs). L-Glu-gated ion channels, more commonly known as (iGluRs), are the primary mediators of fast excitatory synaptic transmission between neurons in the mammalian CNS. Evidence suggests that the iGluR receptor systems are involved in a number of pathophysiology conditions. This chapter will detail the current state of knowledge regarding the involvement of iGluRs in neurodegenerative and other disorders.

#### Keywords

Alzheimer's disease • Excitotoxicity • Ionotropic glutamate receptor • Neurodegeneration • Parkinson's disease

#### List of Abbreviations

AD	Alzheimer's disease
AMPA	Alpha-amino-3-hydroxy-5-methyl-4-isoxazole propionate
AMPAR	AMPA receptor
CNS	Central nervous system
EAA	Excitatory amino acid
iGluR	Ionotropic glutamate receptor
Kainate	(2S,3S,4S)-3-(carboxymethyl)-4-prop-1-en-2-ylpyrrolidine-2-
	carboxylic acid
L-Glu	L-glutamate
NMDA	(2R)-2-(methylamino)-butanedioic acid
NMDAR	NMDA receptor

## 1 History of Central Nervous System Disorders

Disorders of the central nervous system (CNS) have been a part of recorded history since ancient times. The Greek Athenian philosopher Socrates (469–399 BC) is

believed to have experienced neuropsychological disturbances in the form of "demonic orders" or what are now medically referred to as auditory command hallucinations. Similarly, the Greek historian Aristotle (384–322 BC) who was said to have suffered from epilepsy was the first to propose that all mental illness was caused by physical problems of the body (see Traicté de l'Epilepsie by Jean Taxol, 1602). Texts also allude to various other notable historical figures being afflicted with neurological disorders including Julius Caesar (epilepsy), Mozart (manic depression), and Joan of Arc (schizophrenia). Neurological disorders have not only been part of the human condition since the beginning of civilization but also played pivotal roles in human history.

## 2 Multifactorial Etiology of CNS Disorders

Disorders of the CNS are complex disease states with more than 600 disorders currently recognized. Of all CNS disorders, neurodegenerative diseases are the most prevalent. Etymologically, neurodegeneration is made up from prefix "neuro-," which refers to cells of neuronal origin, and "degeneration," which in the context of an organ or tissue system refers to the process of losing function or structure. Clinically, neurodegenerative diseases comprise a diverse assemblage of disorders that present with heterogeneous pathological and clinical expression patterns. These diseases are typically characterized by nervous system dysfunction resulting from the progressive loss of neuron structure or function often with accompanying neuronal cell death. Cell death primarily affects subsets of neurons in specific functional systems and may involve either the entire neuron (neuronopathy) or be restricted to the axon (axonopathy). While the etiology of neurodegenerative diseases is often unknown, a variety of factors have been identified that determine an individual's susceptibility to disease, including an individual's age, race, gender, and genetic makeup. Environmental factors such as exposure to chemicals and infectious agents have also been shown to increase the risk of neurodegenerative diseases. The most prevalent neurodegenerative diseases include Alzheimer's disease and other dementias, Huntington's disease, Parkinson's disease, and amyotrophic lateral sclerosis (ALS or Lou Gehrig's disease).

## 3 Prevalence of Neurodegenerative Disorders

Dementia is the term typically used when referring to a decline in intellectual functioning, including problems with episodic memory, critical reasoning, and objective thinking. Within the general population, the most common disorders that cause neurological impairments include Alzheimer's disease, dementia with Lewy bodies, frontotemporal dementia, and microvascular brain injury. The primary risk factor for dementia is age, with incidence rates doubling for every 5 years after the age of 65. Gender has also been found to affect the incidence of dementia, with woman having a higher risk of developing the condition. Cohort longitudinal

studies estimate that the prevalence rate for all dementias is between 10 and 15 per 1,000 person-years. According to estimates from the World Health Organization, 0.441 % of people worldwide will have dementia by 2015 and that the prevalence will increase to 0.556 % by 2030 (World Health Organization 2012). As of 2010, there were an estimated 35.6 million people worldwide affected by dementia. According to the World Alzheimer Report, released by Alzheimer's Disease International, the prevalence of dementia will double every 20 years to 65.7 million in 2030 and to 115.4 million in 2050 (ASD International 2012). Currently, 58 % of people afflicted with dementia live in developing countries. It is estimated that by 2050 this will rise to 71 %. Alzheimer's disease is the most common form of dementia, accounting for between 50 % and 70 % of all dementia cases.

### 4 Societal Cost of Neurodegenerative Disorders

With a general trend for increasing life expectancies worldwide, neurodegenerative disorders are becoming one of the most costly diseases for societies. In 2009, it was estimated that the approximate cost of dementia worldwide was approximately US\$422 billion yearly. The costs associated with neurodegenerative disorders can be categorized into direct medical costs and indirect costs. Direct medical costs include those associated with long-term care by health-care professionals in addition to the cost of nursing home care. It is estimated that over two-thirds of nursing home residents suffer from some sort of dementia. It is often the case that the primary caregiver for a person suffering from a neurodegenerative disorder is a relative or close friend. This can result in significant indirect costs that include caregiving time and the loss of economic productivity for both the patient and caregiver. It has been estimated that the total costs for an Alzheimer's patient are between \$18,000 and \$77,500 per year in the United States. Dementia caregivers themselves often suffer psychosocial and physical problem which can be directly related to the duties of the caregiver toward the sufferer. It has been estimated that a caregiver may give up to 47 h per week nonwork-related time to the care of a sufferer. It therefore remains that neurodegenerative disorders will continue to represent a major challenge for any society.

## 5 A Brief History of L-Glutamate

In 1866, the German chemist K. Heinrich Ritthausen discovered and identified L-glutamate. However, the role of L-Glu as a nervous system excitant was not fully elucidated until the mid-twentieth century. In 1954, Hayashi observed that direct injection of glutamate into the brain resulted in seizure activities. Similarly, Curtis et al. (Curtis and Eccles 1959; Curtis et al. 1959) showed glutamic acid to be effective for evoking repetitive discharges from interneurons located in the spinal cord of the cat. Subsequent studies by Curtis revealed that application of glutamic acid could induce neuron depolarization with concomitant increases in the firing of

action potentials (Curtis et al. 1959). Concomitant with the presence of high concentrations of glutamate in nervous tissue leads to speculation that L-Glu had an important neurophysiological role. In 1963, a seminal paper by McLennan was published that outlined the criteria required to establish substances as neurotransmitters. In the same year, iontophoretic studies of the effects of L-Glu on neurons in the mammalian cerebral cortex proposed that L-Glu was a putative excitatory neurotransmitter in the CNS (Krnjevic and Phillis 1963). The discovery by Takeuchi and Takeuchi of L-Glu-induced depolarization in muscle at locations corresponding to sites of synaptic innervation added further evidence for the role of L-Glu as a neurotransmitter (Takeuchi and Takeuchi 1963). Since then, numerous experiments have unequivocally determined that L-Glu is the major excitatory neurotransmitter in the vertebrate CNS.

#### 6 Glutamate Receptors

The receptors mediating L-Glu neurotransmission can be classified into two major groups, ionotropic and metabotropic (Fig. 1a). L-Glu-gated ion channels, more commonly known as ionotropic glutamate receptors (iGluRs), are the primary mediators of fast excitatory synaptic transmission between neurons in the mammalian CNS. iGluRs have been typically classified by the synthetic agonists that mimic the effects of L-Glu: (2R)-2-(methylamino)butanedioic acid (N-methyl-D-aspartate; NMDA), 2-amino-3-(5-methyl-3-oxo-1,2-oxazol-4-yl)propanoic acid (AMPA), and (2S.3S.4S)-3-(carboxymethyl)-4-prop-1-en-2-ylpyrrolidine-2-carboxylic acid (kainic acid; kainate) (Fig. 1b). The first cloning of an L-Glu receptor was by Gunderson (1984) who observed that messenger RNA (mRNA) from human brain injected into Xenopus oocytes was able to produce both L-Glu- and kainatesensitive channels in addition to voltage-operated channels. In 1989, Hollmann screened a rat brain cDNA library for kainate-gated ion channels that were subsequently expressed in *Xenopus* oocytes (Hollmann et al. 1989). The resulting cDNA clone was found to encode a single protein of 99,800 relative molecular mass that formed a functional ion channel that possessed both pharmacological and electrophysiological properties of the kainate subtype of L-Glu receptor. Based on its sensitivity to kainate, the receptor was named GluR-K1. However subsequent binding studies of this channel revealed that the receptor was in fact an AMPA receptor protein based on its non-desensitizing responses to kainate. The GluR-K1 receptor was subsequently renamed to GluR1 to reflect this (Sommer et al. 1990).

Over the next few years, additional AMPA receptor family members were cloned including GluR1 (Keinanen et al. 1990), GluR2 (Nakanishi et al. 1990), GluR3 (Bettler et al. 1990; Boulter et al. 1990), and GluR4 (Gallo et al. 1992). In 1990, Bettler et al. cloned a low-affinity kainate-binding subunit they termed GluR5 (Bettler et al. 1990). This receptor subunit displayed only 40–41 % amino acid identity with the other AMPA/kainate receptor subunits known to exist (GluRs1–4). Expression of this clone in Xenopus oocytes resulted in homomeric channels that were only weakly responsive to L-Glu. Investigations determined that

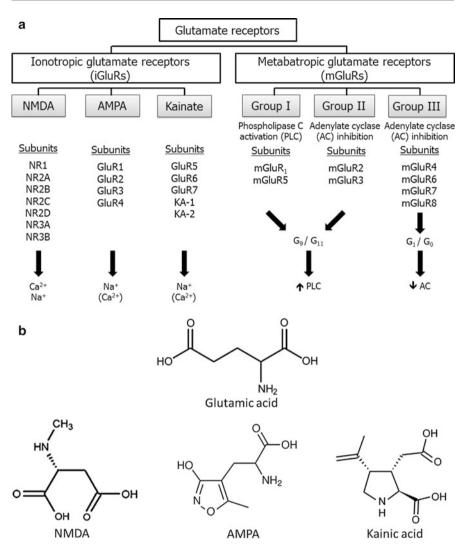


Fig. 1 (a) Family of glutamate receptors. (b) Structures of L-Glu, NMDA, AMPA and Kainate

the sequence similarity of GluR5 was significantly below the 70 % inter-subunit identity that was characteristic of the other AMPA/kainate receptors identified suggesting that GluR5 was in fact a different subtype of L-Glu receptor. This was confirmed in 1992 by Sommer (Sommer et al. 1992) who reported the desensitization of this subunit to kainate. Other kainate-selective receptors were cloned in the early 1990s including GluR6 (Egebjerg et al. 1991), GluR7 (Bettler et al. 1992; Lomeli et al. 1992), KA1 (Werner et al. 1991), and KA2 (Herb et al. 1992; Kamboj et al. 1992). It should be noted that when KA1 and KA2 subunits are expressed either alone or with each other, it does not result in the formation of functional

channels. However, co-expression of KA1 and KA2 with either Glu5 or Glu6 is able to form functional channels albeit with modified properties (Fisher and Mott 2011). The NMDA receptor subunits were first cloned by Moriyoshi et al. (1991), Yamazaki et al. (1992), and Karp et al. (1993) and were termed NR1 (also referred to as NMDAR1). In 1993, Ishii et al. (1993) cloned an additional four subunits, which they termed NR2A-D. Two NR3 subunits have also been cloned termed NR3A and NR3B (Sucher et al. 1995; Das et al. 1998).

Each functional iGluR subtype is formed by the heteromeric assembly of four subunits from a single class, e.g., NMDA subunits can only assemble with other members of the NMDA subfamily. Of the 16 iGluR subunits identified in human, 7 belong to the NMDA class (NR1, NR2A-D, NR3A-B), with the remaining 9 being members of the non-NMDA class, 4 to the AMPA class (GluR1–4) and 5 to the kainate class (GluR5–7 and KA1–2) (Fig. 1a). Two other subunits, known as  $\delta$ 1 and  $\delta$ 2, are termed orphan receptors because while they share high sequence similarity to other iGluR subunits, there is currently no evidence to indicate that they form functional iGluRs (Schmid et al. 2008). Recently, a new nomenclature for ligand-gated ion channels has been proposed by Collingridge et al. (2009) and is currently being reviewed by the nomenclature committee of International Union of Basic and Clinical Pharmacology (IUPHAR) IUPHAR (NC-IUPHAR).

#### 7 NMDA Receptors

The NMDA subtype of iGluRs is involved in activity-dependent forms of synaptic plasticity, synapse consolidation, and excitatory amino acid (EAA)-mediated neuronal toxicity (Hynd et al. 2004a). NMDA receptors (NMDARs) are found on most CNS neurons (Petralia et al. 1994a, b, c) and mediate Ca<sup>2+</sup> influx in response to the coincident binding of L-Glu and glycine (Constantine-Paton 1990; Bliss and Collingridge 1993; Seeburg 1993).

Functional NMDARs exist as tetrameric or pentameric complexes comprising proteins from three families of homologous subunits, designated NR1, NR2A-D, and NR3A-B (Monyer et al. 1992; Ishii et al. 1993; Dingledine et al. 1999). The presence of the NR3 subunits serves to have an inhibitory effect on NMDAR activity. The NR1 subunit functions to bind glycine, whereas the NR2 subunits bind L-Glu. For both the NR1 and NR2 mRNA transcripts and subunit proteins, distinct spatial and temporal expression patterns have been found (Bockers et al. 1994). While the NR1 subunit is expressed ubiquitously in the brain, the NR2 subunits show marked regional and developmental variations (Watanabe et al. 1992; Akazawa et al. 1994; Monyer et al. 1994). The function and modulation of NMDAR channels is both spatially and temporally regulated (Bockers et al. 1994; Rigby et al. 1996). The NR1 mRNA undergoes alternate splicing of three exons (4, 20, and 21) to give eight functionally distinct subunit variants (Sugihara et al. 1992; Durand et al. 1993; Hollmann et al. 1993). Exon 4 encodes a 21-amino acid amino-terminal cassette (designated "N1"), while exons 20 and 21 encode two consecutive carboxy-terminal 37- ("C1") and 38-amino acid ("C2") cassettes

(Zimmer et al. 1995). Both the gene sequence and the intron/exon organization of the human NR1 gene show a high degree of homology to the rat gene, with exception of the deletion of rat exon #3 (Hollmann et al. 1993). In vivo, NMDARs are regulated by a variety of agents including glycine, Mg<sup>2+</sup>, Zn<sup>2+</sup> and H<sup>+</sup> ions, and polvamines. Alternative splicing of the N1 cassette has been shown to modulate both the ligand affinity and channel characteristics of NMDARs. Insertion of the N1 cassette results in the formation of receptors with decreased sensitivity to proton inhibition, increased current amplitudes, and altered responses to protein kinase C activation (Durand et al. 1993; Traynelis et al. 1995). Deletion of the cassette potentiates responses to the polyamines spermine, spermidine, and Zn<sup>2+</sup> (Zhang et al. 1994; Traynelis et al. 1995). In contrast deletion of the N1 cassette has been shown to potentiate the response to polyamines in the presence of saturating concentrations of glycine (Durand et al. 1993) in addition to low-micromolar  $Zn^{2+}$  concentrations (Hollmann et al. 1993). In contrast, both N-terminal variants are inhibited by high-micromolar concentrations of Zn<sup>2+</sup> and at low glycine concentrations. Potentiation of NMDARs occurs dose dependently and involves an increase in both peak channel open probability and peak channel current duration (Araneda et al. 1993).

## 8 AMPA Receptors

The AMPA type of iGluRs (AMPARs) are believed to be the most common type of neurotransmitter receptor in the CNS. AMPARs are multimeric ligand-gated ion channels that like NMDARs play crucial roles in synaptic transmission and plasticity (Ozawa et al. 1998). AMPARs are heteromeric and are composed from four different subunit types, designated as GluR1, GluR2, GluR3, and GluR4 (previously referred to as GluRA-D, respectively) (Collingridge et al. 2009). Each of the subunits is encoded for by a separate gene. Native AMPARs typically occur in a symmetric "dimer of dimers" configuration, consisting of GluR2 and either GluR1, GluR3, or GluR4. Dimerization of the subunits is initiated in the endoplasmic reticulum and occurs via interaction of a conserved N-terminal domain consisting of the amino acids leucine, isoleucine, and valine-binding protein motif. The association of different GluR subunits gives rise to receptors with different channel properties and ionic conductances (Derkach et al. 2007). Each GluR subunit consists of approximately 900 amino acids with each exhibiting 65–75 % sequence homology when compared to the other subunits (Hollmann and Heinemann 1994).

Similar to other iGluRs, the GluR subunits possess a structure consisting of an extracellular N-terminal domain, an intracellular C-terminal domain, and three membrane-spanning hydrophobic domains termed M1, M3, and M4. In addition, one membrane reentrant loop (M2) is also present (Bennett and Dingledine 1995). Within each subunit, the ligand-binding domain is made up from an N-terminal region designated as S1 and S2. However, AMPAR subunits show the greatest difference in their C-terminal sequence, which determines how each subunit interacts with cellular scaffolding proteins. Each AMPAR has four agonist-binding

sites, each of which is formed by the N-terminal region and the extracellular loops located between TM's III and IV. The binding of an agonist (such as L-Glu) results in a structural rearrangement of these two loops, such that each is moved toward each other, resulting in opening of the channel pore. Ion flow through AMPARs only occurs upon coincident binding of two agonists with increases in current flow occurring following binding of two more agonists at the unoccupied sites. Due to the rapid nature for both AMPAR desensitization and resensitization kinetics, it has been suggested that they are the primary mediators for fast excitatory neurotransmission in the CNS. Long-term potentiation (LTP) via AMPARs occurs by both an increase in the number of AMPARs at the postsynaptic surface and increases in the single channel conductance of the receptor channel (Lee and Kirkwood 2011).

Native AMPAR subunits exist as two splice variants, known as the "flip" and "flop" types. Alternative splicing is based on the insertion/deletion of a 38-amino acid sequence located at the C-terminal end of an extracellular loop located between transmembrane segments III and IV (Seeburg et al. 1998). The resulting AMPARs show altered desensitization kinetics including the speed of receptor desensitization, the speed at which the receptor is subsequently resensitized in addition to the rate at which the AMPAR channel closes (Frandsen and Schousboe 2003). The flip variant has been shown to occur in the prenatal CNS, while the flop variant is the form found in the adult CNS.

AMPARs are permeable to the monovalent cations, Na<sup>+</sup> and K<sup>+</sup>. However, in contrast to NMDARs, AMPARs are typically impermeable to the divalent cation, calcium. Ca<sup>2+</sup> permeability of AMPARs is controlled by the GluR2 subunit. Posttranscriptional editing of the GluR2 mRNA results in a single amino acid change from the uncharged amino acid glutamine (Q) to the positively charged arginine (R). This substitution occurs in transmembrane region II, known as the Q/R editing site. AMPARs containing glutamine, GluR2(Q), are permeable to Ca<sup>2+</sup>, while receptors containing arginine, GluR2(R), are not due to the unfavorable energetics required for calcium to enter past the positive-charged amino acid. In the adult CNS, the majority of GluR2 subunits exist as the GluR2(R) form, resulting in Ca<sup>2+</sup>-impermeable AMPARs. This impermeability is believed to protect against neuronal excitotoxicity. The GluR2 subunit has also been shown to modulation of the receptor by polyamines (Stromgaard and Mellor 2004). AMPARs lacking a GluR2 subunit are capable of being blocked in a voltagedependent manner, preventing the passage of positive ions (K<sup>+</sup>, Na<sup>+</sup>) through the channel pore.

The phosphorylation of AMPAR subunits has been shown to alter both synaptic transmission and plasticity in the CNS. Phosphorylation of AMPARs at the intracellular carboxy-terminus has been well established. Serine/threonine phosphorylation sites have been identified on all four AMPA subunits and serve to regulate channel localization, open probability, and ionic conductance (Santos et al. 2009). The GluR1 subunit has four phosphorylation sites which are located at serine 818 (S818), S831, and S845 and threonine 840 (T840) (Lee et al. 2007). Phosphorylation of S818 by protein kinase C (PKC) has been shown to play a crucial role in

long-term potentiation (LTP) (Boehm et al. 2006). Similarly, during LTP, S831 is phosphorylated resulting in the trafficking of GluA1-containing AMPARs to the synapse (Lee et al. 2000). The phosphorylation of S845 by protein kinase A has been shown to regulate channel open probability (Castellani et al. 2005). In contrast, the T840 site has been implicated in long-term depression (LTD) (Lee et al. 2007).

## 9 Kainate Receptors

The kainate types of iGluRs are formed from five distinct subunit types – GluR5 (GluK1), GluR6 (GluK2), GluR7 (GluK3), KA1 (GluK4), and KA2 (GluK5). Similar to both the NMDA and AMPA receptors, kainate receptor subunits can associate in multiple ways to form tetrameric receptors (Contractor et al. 2011). In vivo, the GluR5–7 (GluK1–3) subunits associate to form both homomeric receptors (i.e., receptors that are composed only of one type of GluR subunit) and heteromeric receptors (i.e., receptors whose subunit composition consists of different GluR subunits). In contrast, both the KA1 and KA2 (GluK4 and GluK5) subunits can only form functional receptors when combined with one of the GluR5–7 (GluK1–3) subunits.

Similar to other iGluRs, kainate receptors share a high degree of structural similarity to other subunits of the iGluR family. Receptors consist of an extracellular N-terminal domain that is involved in receptor assembly that, together with a loop between TMIII and TMIV, forms the ligand-binding domain (S1) and a reentrant loop (TMII) that forms the lining of the pore region of the ion channel. Entry of the TMII segment occurs only partially into the plasma membrane before dipping back to the cytoplasm. This segment has been termed the "p loop" and determines the calcium permeability of the receptor. GluRs 5-7 (GluK1-3) have been shown to occur in numerous splice variants and to undergo both RNA editing, giving rise to a variety of receptor types with differing pharmacological and functional properties (Jaskolski et al. 2004). The ion channels formed by kainate receptors are permeable to  $K^+$  and  $Na^+$ , but typically impermeable to  $Ca^{2+}$ . Similar to AMPARs, the ionic conductance of kainate receptors is approximately 20pS; however, their channel kinetics are significantly shorter in duration when compared to AMPARs. Association of receptors with the protein Neto1 has been shown to slow the activation and deactivation kinetics of synaptic responses (Lerma 2011). By modulating the likelihood of depolarization of the postsynaptic neuron, kainate receptors can directly affect the amount of neurotransmitter that is released into the synaptic cleft (Huettner 2003).

Similar to both NMDA and AMPA receptors, kainate receptors are widely distributed throughout the brain. Based on this distribution it has been suggested that rather than being direct mediators of synaptic transmission, the receptors may instead play a role in modulating synaptic plasticity. The spatial distribution for the GluR5–7 subunits is found in somewhat higher levels in the CA3 region of the hippocampus, the striatum, and inner layers of the cerebral cortex. In contrast, both

the KA1 and KA2 subunits are more spatially restricted, with KA1 present only in the hippocampus (Darstein et al. 2003) while KA2 subunits found in the striatum and the inner/outer layers of the cortex (Gallyas et al. 2003). Kainate receptors are the least studied amongst those in the iGluR family, which is largely due to the lack of selective agonists and antagonists for this subtype. Kainate receptors have been implicated in several disease states from clinical data, autopsy brain studies, and animal models including knockout mice (Swanson 2009; Matute 2011). However, due to the crossover activity of many antagonists with AMPARs, it has been difficult to specifically determine the contribution of kainate receptor block to specific neurodegenerative disorders. NBQX, which has shown improvement of akinesia in rat models of Parkinson's disease, is an antagonist at both AMPA and kainate receptors, but is more potent at AMPA receptors. Thus, for the purpose of this chapter, mainly the effects of NMDA and AMPA receptors will be discussed.

#### 10 L-Glu-Mediated Neuronal Excitotoxicity

The first report of the neurotoxic effects of L-Glu was published by Hayashi in 1954 following the observation that direct application of L-Glu was sufficient to induce seizure activity. Subsequent studies by Lucas and Newhouse (1957) showed that subcutaneous injection of L-Glu into postnatal mice resulted in degeneration of neurons in the retina. In 1969 Olney coined the term "excitotoxicity" when he showed that L-Glu-mediated cell death was not limited to the retina but could also occur in the CNS. Application of L-Glu could result in the selective death of postsynaptic neurons and that glutamate agonists could act as potent neurotoxins. Based on the ability of L-Glu to activate iGluRs and that L-Glu antagonists were able to mediate the resulting neurotoxicity, only the term "excitotoxicity" was coined, a portmanteau of excitatory and neurotoxicity. Excitotoxicity is thus defined as the pathological processes by which neurons are either damaged or killed as a result of excessive stimulation by excitatory amino acids (EAAs).

Neuronal excitotoxicity occurs when iGluRs are repetitively activated by endogenous EEAs. As the primary mediator of excitatory neurotransmission in the CNS, L-Glu is considered to be the major endogenous excitotoxin in the brain. Within the brain, L-Glu is synthesized from L-glutamine and Krebs cycle intermediates. When viewed as a complete organ, the concentration of *intracellular* L-Glu is between 5 and 15 mmol/kg wet weight, which far exceeds the *extracellular* L-Glu concentration, which is normally in the range of 0.2–5 mmol/L (Erecinska and Silver 1990). Within presynaptic neurons, L-Glu is stored in vesicles whose contents are subsequently released into the synaptic cleft via a Ca<sup>2+</sup>-dependent mechanism. Once in the synaptic cleft, L-Glu binds to – and activates – postsynaptic iGluRs. Under normal physiological conditions, L-Glu is present in the synaptic cleft at a concentration of 0.6  $\mu$ M (Bouvier et al. 1992). During synaptic transmission, the concentration of L-Glu in the synaptic cleft is increased up to 1 mM, which is then rapidly returned to normal over a millisecond time course (Clements et al. 1992). The excitatory action of L-Glu is terminated by its removal from the synaptic cleft. To date, two primary mechanisms for synaptic L-Glu removal have been identified. The first involves the reuptake of L-Glu into the presynaptic neuron, while the second occurs via a glial cell-based high-affinity reuptake system (Castagna et al. 1997; Kanai et al. 1994). As a result of these sodium-dependent transport systems, a 5,000-10,000-fold concentration gradient of L-Glu exists between the intracellular and extracellular neuronal environments (Ferrarese et al. 1993).

The overactivation of iGluRs by L-Glu results in excessive amounts of  $Ca^{2+}$  entering the postsynaptic neuron. This influx of calcium can result in the sequential activation of a number of degradative enzymatic, nucleolytic, and proteolytic pathways (Zundorf and Reiser 2011). As a result, cellular DNA, RNA, and proteins begin to be degraded along with dismantling of the cytoskeleton and cell membrane. With continued L-Glu stimulation and concomitant calcium ion influx, the neuron undergoes cellular degeneration. Excitotoxic theories suggest that the cellular mechanisms normally involved in excitatory neurotransmission can be transformed into mechanisms of neuronal cell destruction (Olney et al. 1986; Whetsell and Shapira 1993). EAA neurotoxicity in animal models has been found to be cellularly selective, with distinctive degenerative patterns produced by different EAAs (Coyle et al. 1981).

## 11 Excitotoxicity and Neurodegenerative Disorders

Excitotoxicity is thought to be a major contributing factor to a variety of degenerative disorders, including spinal cord injury and hearing loss (ototoxicity), and in neurodegenerative diseases of the central nervous system (CNS). Excessive activation of iGluRs has been implicated in a variety of neurodegenerative diseases including traumatic brain injury (Luo et al. 2011), amyotrophic lateral sclerosis (ALS) (Grosskreutz et al. 2010), multiple sclerosis (Frigo et al. 2012), epilepsy (Czuczwar 2000), hypoxia-ischemia, and chronic neurodegenerative disorders such as Alzheimer's (Woods and Padmanabhan 2012), Parkinson's (Surmeier et al. 2010), and Huntington's diseases (Raymond et al. 2011).

## 12 Involvement of iGluRs in Alzheimer's Disease

Alzheimer's disease (AD) is the most frequent cause of cognitive deficit in the aged population with the prevalence of probable AD is 3 % in subjects aged 65–74, 19 % in those aged 75–84, and 47 % in those over 85. AD represents the fourth most common cause of death in Western countries (Forsyth and Ritzline 1998). With the continuing increase in the world's aged population, projections indicate that by 2050 over 115 million people will suffer from AD (World Health Organization 2012). While normal cognitive losses do occur as part of the normal aging process, the cognitive impairments observed in dementia patients are both qualitatively and

quantitatively more severe (Roses 1998a, b). Following postmortem histopathological examination,  $\sim$ 75 % of patients that presented with a clinical diagnosis of dementia are confirmed to have AD (Evans 1989).

The impairments in memory and cognition observed in AD can be correlated to the neuropathological features of the disease, including neuronal loss and plaque and tangle formation in both the cortex and hippocampus (Hyman et al. 1986; Albin and Greenamyre 1992; Bobinski et al. 1998). While neuronal loss is generally restricted to the pyramidal cells in layers III and IV, damage to L-Glu-innervated neurons is also observed. In AD, neuronal cell death is generally restricted to neuronal cell bodies and dendrites, with axons, terminal boutons, and glial, endothelial, and ependymal cells relatively spared. There is mounting evidence that the region-selective pattern of neuronal cell destruction may occur as a consequence of EAA dysfunction, resulting in the cognitive symptomatology observed. The neurodegeneration observed in AD may therefore arise from both the regional and cellular distribution of an EAA defect, with the observed pathology resulting from the heterogeneous distribution of EAA receptor subtype(s), including NMDA, AMPA, and the kainate receptor types (Hynd et al. 2004a).

#### 12.1 AMPA Receptors in AD

Data from both experimental and clinical trials have suggested that modulation of AMPA receptors may be therapeutically effective in the treatment of neurodegenerative diseases. Reduced levels of the AMPA receptor subunits GluR1, GluR2, and GluR2/3 have been found in the pathologically susceptible areas of the AD brain, including the CA1 region of the hippocampus (Armstrong et al. 1994; Aronica et al. 1998; Garcia-Ladona et al. 1994; Ikonomovic et al. 1997; Wakabayashi et al. 1994). In AD patients, reductions in AMPAR expression can be correlated to disease progression and disease severity (Carter et al. 2004). Within the subiculum, a reduction in GluR2 and GluR2/3 protein levels was found to correlate with Braak stages I–II and stages III–IV, respectively. While in Braak stages V–VI, both GluR2 and GluR2/3 protein levels were similar to those of Braak stages I-II. In contrast, despite reduced subiculum GluR1 protein levels in patients with severe AD, a direct correlation between Braak staging and protein loss throughout all stages of the disease has not been demonstrated (Armstrong et al. 1994; Schaeffer and Gattaz 2008; Yasuda et al. 1995). Within the pathologically spared regions, hippocampal regions, CA2 and CA3, and the dentate gyrus, AMPAR subunits show relatively little change across all Braak stages.

AD is typically characterized by the presence of amyloid plaques and intracellular neurofibrillary tangles. Amyloid is the general term given for peptides produced by the cleavage of amyloid precursor protein (APP). The amyloid beta (A $\beta$ ) peptide is the primary component of amyloid plaques, which are thought to be triggered by the self-aggregation of A $\beta$  into oligomers. It is the near ubiquitous presence of both amyloid plaques and neurofibrillary tangles that is thought to contribute to the degeneration of neurons and subsequently to the progressive decline in cognitive function observed in AD (Shankar and Walsh 2009). Loss of GluR2/3 immunoreactivity has been shown to precede the formation of neurofibrillary tangles in both the hippocampus and entorhinal cortex in the brains of AD sufferers (Ikonomovic et al. 1997). A reduction in the binding of AMPA to receptors has also been observed in AD cases when analyzed by quantitative autoradiography (Dewar et al. 1991). The trafficking of AMPARs has also been shown to be disrupted by soluble A $\beta$  peptides (Shankar et al. 2007).

 $Ca^{2+}/calmodulin-dependent protein kinases II (CaMKII) are serine/threonine$ specific protein kinases involved in signaling cascades that are important in $mediating long-term potentiation (LTP). A<math>\beta$  has been found to impair AMPAR receptor trafficking and function by reducing CaMKII synaptic distribution (Gu et al. 2009). By competition with the proteolytic maturation of brain-derived neurotrophic factor (BDNF), A $\beta$  can also interfere with synaptic potentiation by reducing AMPAR trafficking (Zheng et al. 2010). Similarly, overexpression of APP in cultured hippocampal neurons has been shown to result in reductions in the receptors and cytoskeletal elements at synapses. In PSD-95 knockout mice, reduced AMPAR receptor activity is observed. The loss of PSD-95 may contribute to AMPAR-dependent LTP deficits observed in AD. Significantly decreased levels of PSD-95 have been found in the AD brain (Sultana et al. 2010). Altered AMPA receptor currents that correlate with the reduced GluR2/3 immunoreactivity precede NFT deposition and synapse loss (Ikonomovic et al. 1997).

#### 12.2 NMDA Receptors in AD

Many neurodegenerative disorders, including AD, involve abnormalities in synaptic structure and function. NMDARs show altered expression in a variety of neurodegenerative diseases, including AD and Parkinson's disease (PD). Immunohistochemical analysis of the distribution of NMDAR subunits in AD has to date yielded conflicting results. In a study of the AD patients with mild/modest pathology, staining for the NMDAR in the hippocampus was found to be indistinguishable from that of controls. However, in cases with severe AD changes, a more complex staining pattern was observed (Ikonomovic et al. 1999). Furthermore, quantitative autoradiography has revealed substantially higher binding to NMDARs in the striatum and nucleus accumbens in AD patients (Ulas et al. 1994). A follow-up study however showed significantly lower NR1 mRNA levels in layer III of the entorhinal cortex in AD brains using in situ hybridization. Similar reductions were also observed in layers II and IV-VI of the entorhinal cortex and in granule cells of the dentate gyrus. However, no differences in mRNA expression were observed in the CA1 region of the hippocampus or the subiculum (Ulas and Cotman, 1997). Locally lower NMDAR levels have been reported in the AD hippocampus, although no reductions were noted in either the occipital cortex or caudate (Sze et al. 2001). The basis of these discrepancies may be a result of alternative splicing of the NMDA NR1 mRNA that yields distinct protein subunit variants.

In the AD brain, regional variations in the expression of NMDA NR1 splice variants have been shown to occur between areas that are pathologically susceptible and those that remain relatively spared (Hynd et al. 2004b, 2001). The N1 (exon 4) cassette of the NR1 mRNA is particularly important in the modulation of the mature receptor. Exon 4 encodes a 21-amino acid segment in the N-terminal domain (Durand et al. 1993). Insertion/deletion of the N1 cassette yields NMDAR splice variants (designated the NR1<sub>1XX</sub> and NR1<sub>0XX</sub> variants, respectively) that are subject to distinct modulation by endogenous molecules, including the polyamines spermidine, spermine, and putrescine. Polyamines are a group of organic polycations that are protonated at physiological pH. Polyamines are present in all nucleated cells and occur in the mammalian brain in micromolar concentrations (Sarhan and Seiler 1989). They modulate various cellular functions, including cellular growth and differentiation (Tabor and Tabor 1984), regulation of gene expression (Celano et al. 1989), protein synthesis and phosphorylation (Lenzen et al. 1986), and programmed cell death (Ha et al. 1997). The action of polyamines on the NMDAR complex consists of both stimulatory and inhibitory components (Reynolds and Miller 1989; Williams et al. 1989; Rock and Macdonald 1995). While polyamines do not directly activate the NMDAR, they act to potentiate or inhibit L-Glu-mediated responses in the presence of glycine. Both NR10XX and NR11XX variants are inhibited by high-micromolar Zn<sup>2+</sup> concentrations and potentiated by polyamines at low glycine concentrations (Zhang et al. 1994). However, only NR1<sub>0XX</sub> variants are potentiated by polyamines at saturating glycine concentrations (Durand et al. 1993). The concentration of polyamines in the brain shows marked changes in response to pathological state (Paschen 1992). In particular, levels of brain polyamines are altered in AD (Morrison et al. 1995; Morrison and Kish 1995). In AD cases, mean levels of spermidine, the immediate biosynthetic precursor to spermine, are markedly increased compared to controls in the temporal cortex of patients. Conversely, putrescine levels are decreased in the same region in AD cases. In one study, the ratio of NR1 1XX:0XX was found to be significantly reduced in regions of the AD brain most susceptible to pathological damage (Hynd et al. 2001), while levels of the NR1<sub>1XX</sub> mRNA were found to be lower in cingulate gyrus, the hippocampal formation, and superior temporal gyrus in AD cases. Conversely, NR11XX levels were unchanged in both the motor and occipital cortices in AD cases and controls. These results suggest that, at least in the pathologically susceptible temporal cortex of AD patients, stimulation of the NMDAR (as a consequence of elevated spermidine levels) may be increased and its antagonism (through lowered putrescine levels) would be decreased. The selective vulnerability of neurons in AD may be attributed, in part to local variations in the composition and hence function of the NMDA NR1 receptor.

The co-expression of NR1 variants with an NR2(A–D) subunit results in heteromeric channels with distinct electrophysiological and pharmacological profiles. Significantly lower levels of both the NR2A and NR2B mRNAs were found in susceptible brain regions, including the hippocampus, temporal, and cingulate cortices (Hynd et al. 2004b).

#### 13 Involvement of iGluRs in Parkinson's Disease

Parkinson's disease (PD) is the second most common neurodegenerative disorder in the world after Alzheimer's disease, affecting about 1 % of adults over 60 years of age. PD is a disease which is more common in the elderly and prevalence rises from 1 % in those over 60 years of age to 4 % of the population over 80 (de Lau and Breteler 2006). Although the mean age of onset is around 60 years, 5–10 % of cases are diagnosed as young onset PD between the ages of 20 and 50 (Samii et al. 2004).

The hallmark of PD pathology is loss of dopaminergic neurons in the substantia nigra pars compacta of the midbrain, which manifests itself with symptoms such as bradykinesia, muscle rigidity, and tremor. Bradykinesia involves delayed voluntary movements, difficulty in undertaking rapid motor tasks and multitasking, and also decreased facial movements. Rigidity affects the patient's posture because of an increase in tone throughout all movements. The tremor or shaking primarily affects the hands, arms, legs, jaw, and face. As these symptoms become more pronounced, patients may have difficulty walking, talking, or completing other simple tasks. The first symptoms appear after a large proportion of the dopaminergic neurons are lost. The severity of how PD affects different people varies greatly and there is no way to predict the symptoms. Although some people become severely disabled, others experience only minor motor dysfunctions.

Current established therapies focus primarily on alleviating the motor symptoms using dopamine replacement therapy with levodopa (L-DOPA) (Calabresi et al. 2010; Wu and Frucht 2005), which is a dopamine precursor. However, long-term complications after continuous use of levodopa involve disabling fluctuations and dyskinesias (Calabresi et al. 2010).

However, the chronic and progressive neurodegeneration in PD also affects non-dopaminergic pathways, which causes nonmotor symptoms, such as autonomic dysfunction, cognitive and psychiatric changes, sensory symptoms, and sleep disturbances (Samii et al. 2004). The striatum is the principal input structure of the basal ganglia, where excitatory glutamatergic input and dopaminergic input convey and interact. The depletion of nigrostriatal dopaminergic neurons leads to overactivity of the glutamatergic pathways within the striatum and output nuclei. The involvement of the glutamatergic system in the pathogenesis and symptomatology of PD is well established (Blandini et al. 1996; Chase and Oh 2000; Lange et al. 1997; Starr 1995), and the iGluRs are of interest as potential targets for anti-PD treatment therapies.

#### 13.1 NMDA Receptors in PD

NMDARs are widely expressed in the CNS, including all of the structures of the basal ganglia, but with different subunit compositions. The abundance of NMDA receptors in the postsynaptic density of glutamatergic synapses suggests an important role in rapid response to neurotransmitter release and modulation of glutamate excitation in both physiological and pathological conditions, including PD.

Due to the presumed involvement of NMDARs in PD, NMDA receptor antagonists have therefore been of major interest for the treatment of PD in recent years, but results from both experimental studies and human trials have given mixed results. In one study by Calon (Calon et al. 2003), tissue obtained at autopsy showed increased binding of 3H-Ro 25–6981 (an NR1/NR2B-selective antagonist) in the lateral putamen when compared to controls, while binding of 3H-CGP39653 (an NR1/NR2A-selective antagonist) was unchanged in PD patients with levodopa-induced motor complications. Another study reported a decrease in the abundance of NR1/NR2B subunits in the striatum of a parkinsonian model of the rat. In contrast, no changes in the NR2A subunit were found (Dunah et al. 2000). An in vivo study used positron emission tomography to compare NMDA receptor activation in PD patients with and without levodopa-induced dyskinesias showed an increased activity in patients with dyskinesias compared to those without dyskinesias (Ahmed et al. 2011).

The noncompetitive low-affinity NMDA receptor antagonist amantadine in combination with levodopa or as monotherapy has showed limited antiparkinsonian effects and reduction in dyskinesia in PD patients (Crosby et al. 2003). Remacemide, another low-affinity NMDA receptor antagonist, showed promising antiparkinsonian effects in rodent and primate models of PD (Greenamyre et al. 1994), but when tested in 279 PD patients with motor symptoms and treated with levodopa, it showed only modest improvements although not significant (Shoulson et al. 2001). Therapeutic levels of MK-801, a noncompetitive NMDA receptor antagonist, suppress dyskinesia but also worsen parkinsonism such as sensorimotor deficits in the 6-hydroxydopamine (6-OHDA) rat model (Paquette et al. 2010). Memantine, an NMDA receptor antagonist, showed promise in treating PD patients with motor symptoms taking levodopa (Rabey et al. 1992), but in a larger study with PD patients with dementia, there was no effect on motor symptoms (Aarsland et al. 2009). However, there was an overall improvement in cognitive functions. More recent studies have focused on subtype-selective NMDA receptor antagonists, primarily those selective for the NR2B subunit, with expectation to achieve antiparkinsonian effects without the adverse effects seen from nonspecific antagonists. However the effects of NR2B-selective NMDA receptor antagonists as PD therapeutics have been mixed. The NR1/NR2B-specific NMDA receptor antagonist CP-101,606 showed limited effect on parkinsonian symptoms when administered as monotherapy but showed only a modest potentiation of antiparkinsonian actions of levodopa treatment in PD patients (Nutt et al. 2008) and in a marmoset model of PD (Nash et al. 2004). Another NR2B subtype-selective antagonist, CI-1041, successfully prevented levodopa induced dyskinesias when coadministered, in parkinsonian monkeys (Hadj Tahar et al. 2004). Ro 25-6981 is subtype-specific for active NR2B-containing NMDA receptors and has showed antiparkinsonian effect in both rat and marmoset models of PD, in addition to potentiating levodopa treatment and potentiating dopamine receptor agonists apomorphine, A68930, and quinpirole in rats (Loschmann et al. 2004). Single-dose monotherapy of MK-0657 did not improve motor functions in PD patients, and although the drug was well tolerated, it had adverse effects such as an increase in

blood pressure (Addy et al. 2009). NR2B subtype-selective NMDA receptor antagonists may have potential as therapeutic agents to treat PD, but maybe the mode of action needs to be further investigated so that a drug with fine-tuned properties for best antiparkinsonian effects can be developed.

#### 13.2 AMPA Receptors in PD

AMPARs mediate the vast majority of fast excitatory neurotransmission in the brain and are therefore of interest in regards to the glutamatergic involvement in PD. Increased binding of AMPA in the lateral putamen in PD patients experiencing motor complications, suggests an involvement of AMPA receptors in levodopainduced motor complications (Calon et al. 2003). Several AMPA receptor antagonists have been tested for their antiparkinsonian effect and for their ability to relieve levodopa-induced dyskinesia in rodents (Bibbiani et al. 2005; Megyeri et al. 2007; Stauch Slusher et al. 1995) and monkey models of PD (Bibbiani et al. 2005; Konitsiotis et al. 2000). However, the results are not totally conclusive. The AMPA antagonist GYKI-47261 given in combination with levodopa produced an improvement in dyskinesias; however, significance was not reached (Bibbiani et al. 2005). In contrast, another study showed that the AMPA antagonist LY300164, in combination with various concentrations of levodopa, increased motor activity by up to 86 % and decreased levodopa-induced dyskinesia by up to 40 % (Konitsiotis et al. 2000). GYKI-52466 and GYKI-53405, two noncompetitive AMPA receptor antagonists, exerted only limited effect on dopamine turnover in striatum tissue of rats with experimental Parkinson's when administered alone, but potentiated the effect of levodopa when given in combination (Megyeri et al. 2007). Significant improvements for akinesia were shown in parkinsonian rats after administration of the AMPAR antagonists NBQX, GYKI-52466, and YM90K (Stauch Slusher et al. 1995). Perampanel, another noncompetitive AMPAR antagonist, has been in clinical trials for treatment of PD after showing encouraging results in relieving motor symptoms in animal models. Perampanel was found to be effective in the treatment of refractory partial-onset seizures in three pivotal (Phase 3) clinical trials, although no evidence for improved cognitive functions was found (French et al. 2012).

Although excitotoxicity of the glutamatergic pathways is suggested to play role in PD, stimulatory modulation of AMPA receptors may also have therapeutic potential for the disease. Potentiators of the AMPA receptors, which stimulate synaptic transmission, synaptic plasticity, and cognition, also increase the production of BDNF (Hyman et al. 1991; Lauterborn et al. 2000). BDNF is associated with reduced neurotoxicity and improved neurological signs in cultured dopaminergic neurons (Hyman et al. 1991) and in animal models of Parkinson's disease as shown after LY503430 treatment (Murray et al. 2003). In contrast, the positive AMPA receptor modulator CX516 was reported to cause dyskinesia in a primate model of PD when administered alone or in combination with levodopa (Konitsiotis et al. 2000). The variable results of AMPA potentiators as therapeutics for PD may therefore depend on which class of potentiators they belong to, since they have been suggested to show distinct mode of actions, potencies, and selectivities at different AMPA receptors (O'Neill and Witkin 2007). Further investigations will be required in order to fully determine the extent of AMPAR modulation in PD.

## 14 Traumatic Brain Injury

Traumatic brain injury (TBI), also known as intracranial injury, is a form of acquired primary brain injury that occurs when an external force causes damage to the brain. TBIs can be classified based on the mechanism (i.e., closed or penetrating head injury) and severity of the injury, in addition to the physical location of the injury (e.g., localized or global) (Bruns and Hauser 2003). In addition to the initial primary injury, TBI is often associated with secondary injuries that occur in the hours and days following the primary injury. These injuries result in a cascade of events that serve to alter cerebral blood flow and further contribute to the damage resulting from the primary traumatic insult (Chang et al. 2009). Trauma to the brain can result in vascular damage that results in reduced blood perfusion and may lead to transient ischemia. Ischemia is defined as a restriction in blood supply, with concomitant damage or dysfunction to both cells and tissue. The resulting cerebral hypoxia in which blood flow is reduced to inadequate levels leads to the death of brain tissue or cerebral infarction/ischemic stroke (Bjornsson et al. 2006). TBI typically causes impairment in both cognitive performance and fine motor functions that often persist for months after the initial injury occurs.

## 14.1 NMDA Receptors in TBI

TBI has been shown to cause a marked elevation in the concentrations of extracellular glutamate and aspartate proximal to the trauma site. This results in the excessive activation of NMDARs and neuronal cell death (Bullock 1995; Hovda et al. 1995). Microdialysis studies in humans reveal that there is sustained increases in the concentration of L-Glu lasting between 6 h and several days following TBI (Faden et al. 1989; Choi et al. 1990). Along with this increase, the density of activated NMDARs increases in the hippocampus but decreases in the cortex at the impact site in mice. Interestingly, a study by Kumar (Kumar et al. 2002) using RT-PCR analysis did not show any significant change in the mRNA levels of NR1, NR2A, and NR2B following injury in comparison with controls, suggesting nontranscriptional changes (i.e., protein) occur in the levels of these subunits. Interestingly, protein levels of NR1, NR2A, and NR2B were found to transiently decrease from the time of TBI up until 12 h post-injury before returning to pre-injury levels at 24 h. In contrast, a study by Schumann showed that the expression of the NR1, NR2A, and NR2B subunits were increased in the hippocampus, but decreased in the cortex at the site of the TBI (Schumann et al. 2008). This study revealed that the increase in NMDAR subunit expression in the hippocampus was associated with an increase in NR2B tyrosine phosphorylation. Furthermore, that inhibition of NR2B phosphorylation PP2, a tyrosine kinase inhibitor, was able to restore NMDAR subunit expression to pre-TBI levels. Importantly, inhibition of NR2B phosphorylation was also found to result in significant improvements in long-term recovery of motor functions following TBI.

Administration of NMDA receptor antagonists has been shown to reduce injury severity after TBI (Faden et al. 1989). Pretreatment with either the noncompetitive NMDA antagonist dextrorphan or the competitive NMDA antagonist CGS 19755 was found to limit the release of extracellular L-glutamate following TBI (Panter and Faden 1992). In mice subject to mild TBI, it was observed that cellular viability was reduced in the hippocampus of the mice subjected to trauma while animals preconditioned with 75 mg/kg NMDA no reductions in cellular viability were observed. Similarly, in rats subjected to percussion trauma of the cortex, pretreatment antagonist 3-((+/-)-2-carboxypiperazin-4-yl)-propyl-1with the NMDAR phosphonate (CPP) was successful in mitigating cortical damage (Bernert and Turski 1996). Based on their role in activity-dependent forms of synaptic plasticity (Constantine-Paton 1990; Constantine-Paton et al. 1990) and long-term potentiation (Bliss and Collingridge 1993), NMDARs are believed to play a role in memory and learning following TBI. Mice preconditioned with NMDA and subject to object recognition tasks following TBI were found to be protected against impairments in both long- and short-term memory. In contrast, in mice treated with NMDA, long-term impairments in memory were found when assessed in the step-down inhibitory avoidance test at 7 days following the initial training session (Moojen et al. 2012). Long-term behavioral deficits are often observed in patients following TBI. A study by Hayes (Hayes et al. 1988) showed that pretreatment with phencyclidine, an NMDA antagonist, was able to attenuate the long-term behavioral deficits in the rat produced by TBI. Preconditioning of mice with low-dose concentrations of NMDA has also been used to protect against locomotor deficits observed after TBI. Mice preconditioned with NMDA (75 mg/kg) 24 h prior to TBI induction were found to be protected against all motor deficits revealed by footprint tests. However, this protective effect was not seen when mice were tested in rotarod tasks (Costa et al. 2010). These data suggest that excitotoxicity is mediated by NMDARs with the degree of activation contributing to the pathophysiology of brain injury and ultimately the severity of locomotor dysfunction.

Cerebral hypoperfusion following TBI often occurs as a result of loss of responsiveness to cerebrovascular dilator stimuli via NMDAR-mediated impairment of cerebral hemodynamics. Tissue plasminogen activator is a known activator of NMDARs and is upregulated following TBI. Treatment of mice with a catalytically inactive variant of tissue plasminogen activator was able to prevent impairment of cerebral hemodynamics and decrease excitotoxic neuronal death post-TBI (Armstead et al. 2012). Similarly, the polyamine-site NMDAR antagonist ifenprodil was able to protect against both brain edema and breakdown of the blood–brain barrier after TBI (Dempsey et al. 2000). MK-801, a noncompetitive blocker of the NMDAR, has also been used to treat the neuropathological insults following fluid-percussion brain injury in the rat. In a study by McIntosh, MK-801

treatment significantly attenuated the development of focal brain edema at the site of injury 48 h after TBI (McIntosh et al. 1990). At the molecular level, influx of  $Ca^{2+}$  may not be the only factor contributing to neuronal cell death following elevated concentrations of L-Glu. Recent studies have shown that activation of extrasynaptic NMDARs triggered by L-Glu or hypoxic/ischemic following TBI activates a CREB (cAMP response element-binding protein) pathway. This pathway results in the loss of mitochondrial membrane potential and neuronal apoptosis. These results suggest that pharmacological inhibition of EAA neurotransmission including the use of NMDA antagonists may be used to improve neurological outcome after TBI.

## 14.2 AMPA Receptors in TBI

AMPARs are critical for synaptic plasticity, and a number of recent studies have focused on the role of AMPARs in the physiological and pathophysiological processes following TBI. In vivo, AMPARs are subject to alterations based on subunit composition and receptor trafficking to and from the plasma membrane. A number of recent reports have shown that both the composition and function of synaptic AMPARs change after mechanical injury (Arundine and Tymianski 2003). As a result of these changes, the normally Ca<sup>2+</sup>-impermeable AMPARs can become  $Ca^{2+}$  permeable, thereby affecting calcium signaling and the pathophysiology following TBI. In 2010, Beattie (Beattie et al. 2010) showed that the proinflammatory cytokine tumor necrosis factor (TNF)a, a potent regulator of AMPAR trafficking, can rapidly increase the surface expression of Ca<sup>2+</sup>-permeable AMPARs following TBI. This increase in combination with increased levels of extracellular L-Glu after TBI may play an important role in enhancing excitotoxic cell death after CNS injury. Using an in vitro model of TBI, Spaethling (Spaethling et al. 2008) showed that  $Ca^{2+}$ -permeable AMPARs appear 4 h following injury to cortical neurons. Interestingly, mechanical injury has been shown to modulate AMPAR kinetics via an NMDA receptor-dependent pathway (Goforth et al. 2004). Activation of the NMDAR can lead to CaMKII activation to produce a sustained reduction of AMPA receptor desensitization. Expression of the GluR1 subunit of the AMPAR has also been shown to be significantly decreased in the rat hippocampus following TBI (Kharlamov et al. 2011).

## 15 Involvement of iGluRs in Epilepsy

Epilepsy is a common name for a variety of chronic neurological disorders characterized by seizures resulting from excessive neuronal activity in the brain. The highest incidence occurs in young children and the elderly. These seizures can involve specific systems of the brain (partial seizures) or develop in a restricted area of the brain but then spread to involve multiple cortical and subcortical circuits. In the early 1940s, it was suggested that epileptic seizures occur when the "quantity of the

glutamic acid in human brain surpasses the critical level" after observing that injection of glutamate into the grey matter of the motor cortex of dogs and humans caused epileptic seizures. As the major excitotoxic neurotransmitter in the brain, L-Glu plays a major role in the initiation and spread of seizure activity by acting on iGluRs.

#### 16 AMPA Receptors in Epilepsy

Due to their role in excitatory neurotransmission in the brain, AMPARs have recently become the subject of interest in the pathogenesis of epilepsy. AMPAR antagonists have proven therapeutic potential as antiepileptic drugs in clinical trials. Both talampanel and perampanel, two noncompetitive AMPA receptor antagonists, have proven to reduce seizure frequency in patients with refractory partial seizures (Chappell et al. 2002; Stephen and Brodie 2011). Adverse effects such as dizziness and ataxia were the most common.

#### 17 NMDA Receptors in Epilepsy

At present, the NMDAR is probably the iGluR which has been studied the most as a therapeutic target for treatment of epilepsy. However, despite a large number of studies demonstrating up- or downregulation of particular NMDA receptor subunits' mRNA expression in various animal models of epilepsy, and several NMDA receptor antagonists showing anticonvulsant effect in in vivo animal studies, there is debated evidence for NMDA receptor antagonists as useful treatment for epilepsy in humans (Ghasemi and Schachter 2011). Felbamate, a promiscuous NMDA receptor inhibitor, was approved by the FDA in 1993 but is only used for patients with intractable partial seizures, infantile spasms, or Lennox-Gastaut syndrome, because the potential benefit outweighs the side effects. Remacemide, a low-affinity NMDA receptor blocker, showed improvement as adjunctive therapy in several clinical trials in patients with drug-resistant seizures (Ghasemi and Schachter 2011). Ketamine, an open-channel blocker, has shown both positive and adverse effects in clinical trials. Amantadine, an NMDA receptor blocker, has shown improvements in patients with refractory absence seizures as both mono- and add-on treatment. The results from memantine, an open-channel NMDA receptor blocker, in patients with seizures have been inconclusive despite promising animal studies. In summary, lower doses of NMDA receptor antagonists, such as memantine and remacemide, have shown antiepileptic effects with less severe side effects as adjunct therapy compared with monotherapy (Ghasemi and Schachter 2011).

## 18 Kainate Receptors in Epilepsy

Kainate receptors may play a more modulatory role in fine-tuning excitatory and inhibitory neurotransmission to prevent over excitation, which may lead to epilepsy.

Systemic or intracerebral injection of kainate, an analog of L-Glu, is well known to cause epileptic seizures, originating from the CA3 region of the hippocampus (Vincent and Mulle 2009). Thus, modulating the kainate receptors may have clinical implications in the treatment of epilepsy. Using knockout mice, two independent studies have shown the involvement of the kainate receptor subunit GluK2 in kainate-induced seizures, which are a model for temporal lobe epilepsy (Fisahn et al. 2004; Mulle et al. 1998). Therefore, interfering with GluK2 activity may have potential as antiepileptic drugs. In addition, GluK1 has shown increased expression in the hippocampus of patients suffering from temporal lobe epilepsy (Li et al. 2010), which supports that GluK1 antagonists LY37770 and LY382884 can block seizures in vivo and epileptiform activity in hippocampal slices induced by pilocarpine or electrical stimulation (Barton et al. 2003; Smolders et al. 2002). Topiramate, a GluK1 antagonist, was approved by the Food and Drug Administration (FDA) in 1996 for the treatment of epilepsy both as a mono- and adjunctive therapy.

#### 19 Conclusion

Since its discovery in the 1950s, our understanding of the role that L-Glu plays in both normal synaptic transmission and neurodegenerative disorders has increased dramatically. Numerous experiments have revealed how the molecular mechanism of the human brain can be transformed into implements of neuronal cell destruction. With the advent of gene cloning and transgenic animal models, our understanding of iGluRs and the role they play in neurodegenerative disorders is increasing at an exponential rate. Despite the relative lack of new therapeutics for the treatment of neurodegenerative disorders, the next decade will undoubtedly be one of significant advances for the diagnosis of neurodegenerative diseases and in the development of new and improved treatment regimens for sufferers.

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# Ionotropic Receptors in the Central Nervous System and Neurodegenerative Disease

Nady Braidy, Anne Poljak, Tharusha Jayasena, Gilles J. Guillemin, and Perminder Sachdev

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N. Braidy (⊠) • T. Jayasena

A. Poljak

G.J. Guillemin

St. Vincent's Centre for Applied Medical Research, St. Vincent's Hospital, Sydney, Australia

Neuropharmacology Group, MND and Neurodegenerative Diseases Research Centre, Macquarie University, North Ryde, NSW, Australia e-mail: gilles.guillemin@mq.edu.au

P. Sachdev

Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

e-mail: n.braidy@unsw.edu.au; t.jayasena@student.unsw.edu.au

School of Medical Sciences, Faculty of Medicine and Bioanalytical Mass Spectrometry Facility, University of New South Wales, Sydney, Australia e-mail: a.poljak@unsw.edu.au

Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

Neuropsychiatric Institute, Prince of Wales Hospital, Sydney, Australia e-mail: P.Sachdev@unsw.edu.au

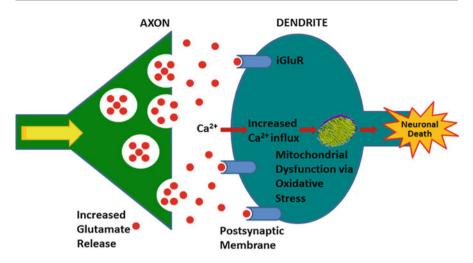
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#### Abstract

Glutamate was identified as the main excitatory neurotransmitter in the mammalian central nervous system (CNS) following the observation in the early 1950s that glutamate can induce seizure activity and excite neurons in the mammalian brain. Over the last two decades, selective ligands, including competitive agonists and antagonists and allosteric modulators, have been developed to further investigate the functional role of glutaminergic receptors. Glutamate released from synapses can activate ligand-gated cation channels at postsynaptic cells to mediate fast postsynaptic potentials. These ion channel-forming ionotropic glutamate receptors (iGluRs) are divided into N-methyl-D-aspartate (NMDA), α-amino-3-hydroxyl-5-methyl-isoxazole-4-prorionate (AMPA), and kainate (KA) receptors. While only 20-30 % of the amino acid sequence is shared among these receptor subtypes, they share similar structural features and their activity is based on specific pharmacological preference. In this chapter, we will describe the structure and composition of iGluRs and infer their pharmacology, with a particular focus on their role in the CNS and their relevance to the pathogenesis of neurodegenerative diseases.

# 1 Introduction

Human neurodegenerative diseases are characterized by extensive loss of neurons, resulting in a number of neurological deficits (Hague, Klaffe et al. 2005). Excitatory amino acids serve as major neurotransmitters in the cerebral cortex and hippocampus. Under normal physiological conditions, excitatory amino acids are essential for the maintenance of normal psychological functions including learning and memory. Dysregulation of the excitatory amino acid system in various CNS diseases is associated with the development of several cognitive pathologies including schizophrenia, delirium, and dementia (Javitt and Zukin 1990). Overactivation of the excitatory amino acid system can also lead to cell death, giving rise to the concept of excitotoxicity (Olney 1978). Excitotoxicity refers to the process of neuronal cell death arising from prolonged exposure to excitatory amino acids with different permeabilities, leading to elevated Ca<sup>2+</sup> influx (Wong, Cai et al. 2002; Berliocchi, Bano et al. 2005). The subsequent  $Ca^{2+}$  overload can stimulate the activation of a number of different enzymes which can degrade proteins, lipids, and nucleic acid (Berliocchi, Bano et al. 2005) (Fig. 1). Apart from glutamate, which is the principal excitatory neurotransmitter in the mammalian CNS, other constitutive amino acids,



**Fig. 1** Schematic representation of glutamate-induced excitotoxicity in neurons. Excitotoxicity due to increased glutamate release from hyperactive neuronal circuits in the brain can activate ionotropic glutamate receptors. Overactivation of these receptors leads to high intracellular calcium influx, disrupts mitochondrial function, and culminates in cell death

such as aspartic acid, and exogenous compounds, such as quinolinic acid and kainate, or of synthetic origin such as NMDA, can induce excitotoxicity (Foster, Collins et al. 1983; Hertz, Dringen et al. 1999).

Neurodegenerative disorders have distinct clinical symptoms and genetic etiologies. While some neurodegenerative conditions are considered "sporadic," due to lack of well-defined genetic associations common to all patients, others do have specific genetic mutations. For instance, the majority of cases of Alzheimer's disease (AD) are sporadic, only around 5 % of these cases are familial forms of autosomaldominant (not sex-linked) inheritance, that usually occurs before the age of 65 (Blennow, de Leon et al. 2006). While these genetic mutations can affect a variety of proteins, sufficient evidence suggests that excitotoxicity represents a convergence point in neuronal degeneration (Dodd 2002; Kawamata and Shimohama 2002). Overactivation of ionotropic glutamate receptors iGluR (Fig. 2) can participate in fast glutaminergic synaptic transmission, leading to impaired calcium homeostasis and increased free radical generation, culminating in programmed cell death (Rae, Moussa et al. 2006).

## 2 Structural Features of Ionotropic Glutamate Receptors

The first transmembrane domain of the iGluR family (Fig. 3) contains a poreforming membrane-residing domain that forms a re-entrant loop entering from and exiting to the cytoplasm (Dingledine and McBain 1994; Dingledine, Borges et al. 1999). A large extracellular loop provides a link between the second and third transmembrane domains, followed by an extracellular carboxy-terminus.

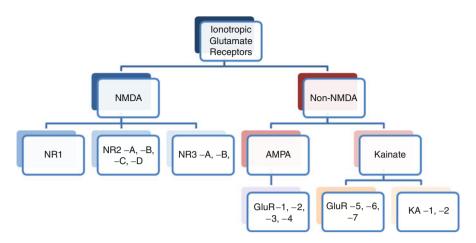
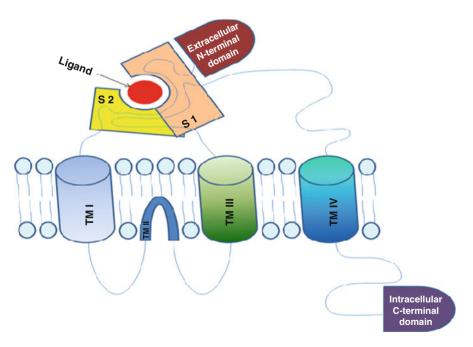


Fig. 2 Family of ionotropic glutamate receptors based on structure and function



**Fig. 3** Schematic diagram of iGluR subunit topology. The extracellular amino-terminal domain is followed by the first transmembrane domain (TM I). The membrane-residing domain (TM II) forms a re-entrant loop. The third and fourth transmembrane domains (TM III and TM IV) are linked by an extracellular loop. The agonist-binding domain is located in a pocket between polypeptide segments S1 and S2

The crystal structure of iGluR ligand binding domains consists of unique polypeptides attached to both the extracellular amino terminus and the extracellular loop between transmembrane domain 3 and 4 (Armstrong, Sun et al. 1998; Mayer and Armstrong 2004).

#### 2.1 NMDA Receptors

NMDA receptors are ligand-gated ion channels that exist as heterotetrameric complexes at the membrane surface. Molecular cloning has identified seven subunits (NR1, NR2A-D, and NR3A and B), each of which represent products of different genes. These amino acid sequences are 18 % (NR1 and NR2), 55 % (NR2A and NR2C), 70 % (NR2A and NR2B), and 25 % (NR3 and NR1, or NR3 and NR2) identical (McBain and Mayer 1994).

Electrophysiological studies have shown that activation of the NMDA receptor requires occupation of two independent glutamate and glycine sites (Clements and Westbrook 1991). Therefore, functional NMDA receptors are composed of the constitutive NR1 subunit and at least one NR2 subunit, or NR1 and both NR2 and NR3 subunits (Mayer and Armstrong 2004). While the exact role of NR3 remains unknown, site-directed mutagenesis has shown that the NR2 subunit contains the glutamate-binding site while the binding site for the co-agonist glycine is carried on the NR1 subunit (Seeburg, Burnashev et al. 1995; Nishi, Hinds et al. 2001; Chatterton, Awobuluyi et al. 2002; Loftis and Janowsky 2003; Matsuda, Fletcher et al. 2003). This glycine-binding domain resembles the clamshell-like architecture reminiscent of the AMPA and kainite subunit-binding pockets (Kew, Koester et al. 2000; Kew and Kemp 2005).

The NR2 and NR3 subunits are differentially expressed in a distinct spatiotemporal pattern (Monyer, Sprengel et al. 1992). In the rodent and human brain, the NR2A and NR2B subunits are predominantly expressed in the forebrain, while the NR2C is expressed in cerebellar granule cells and several nuclei (Monyer, Sprengel et al. 1992; Monyer, Burnashev et al. 1994). However, the expression of the NR2D subunits is confined to the diencephalon and midbrain where it is upregulated significantly during development (Monyer, Sprengel et al. 1992). The NR3 subunit also shares similar spatiotemporal distribution, with the NR3A receptor also highly expressed during development, although expression of the NR3A receptor persists in some neuronal populations during adulthood (Nishi, Hinds et al. 2001). Unlike the NR3A subunit, NR3B is localized in somatic motor neurons in the brainstem and spinal cord. When expressed with NR1 and NR2, the NR3 subunit suppresses NMDA channel unitary conductance and Ca<sup>2+</sup> permeability relative to NR1/NR2containing receptors (Nishi, Hinds et al. 2001; Chatterton, Awobuluyi et al. 2002; Loftis and Janowsky 2003; Matsuda, Fletcher et al. 2003). Moreover, NR3A knockout mice exhibit increased NMDA-mediated currents and increased spine density in cortical neurons (Das, Sasaki et al. 1998). Lee et al. (2011) recently showed that adult and fetal human primary astrocytes express all known NMDA receptor subunits (Lee, Ting et al. 2011). This is in accordance to previous

immunohistochemical studies showing that NR1 and NR2A/NR2B are present in rodent astrocytes (Conti, DeBiasi et al. 1996; Conti, Barbaresi et al. 1999). Native NMDA receptors are therefore composed of NR1 in combination with one or more NR2 subunits (e.g., NR1/NR2A, NR1/NR2B, or NR1/NR2A/NR2B), or alternatively, NR1 may exist in combination with both NR2 and NR3 subunits (e.g., NR1/NR2B/NR3A) (Kew and Kemp 2005).

#### 2.2 AMPA Receptors

Cloning studies have shown that AMPA receptors are composed of four closely related subunits (GluR1, GluR2, GluR3, and GluR4) (Borges and Dingledine 1998). These four subunits appear to be products of separate genes. The polypeptide sequence of each subunit is about 900 amino acids in length and sharing 70 % (GluR1 and 2) and 73 % (GluR2 and 3) homology (Rosenmund, Stern-Bach et al. 1998). Like NMDA receptors, native AMPA receptors are heteromeric in composition and expressed differentially in the brain. GluR2 plays a predictive role in determining the permeability of AMPA receptors to Ca<sup>2+</sup> (Bowie and Mayer 1995). AMPA receptors lacking the GluR2 subunit are Ca<sup>2+</sup> permeable and exhibit evidence of inward channel rectification due to voltage-dependent block of the ion channel by intracellular polyamines (Bowie and Mayer 1995). The molecular property determining the Ca<sup>2+</sup> impermeability of the GluR2 subunit can be attributed to the presence of an arginine (R) positioned at a critical site in the pore loop, M2 domain which is replaced by glutamine (Q) in other AMPA subunits (Seeburg and Hartner 2003).

The diversity of AMPA receptor subunits is due to two genetic processes: RNA editing and alternative splicing. Editing is a posttranscriptional change in at least one base in the pre-mRNA where the codon encoded by the gene differs to the mRNA. In AMPA receptors, editing is restricted to the GluR2 subunit. In the primary transcript of GluR2, glutamine at position 586 in the M2 putative membrane is replaced with an arginine (Dingledine, Borges et al. 1999). The arginine codon is absent in the GluR2 gene and is introduced in the mRNA by dsRNA adenosine deaminase which converts adenosine to inosine which is subsequently read as guanosine. Editing is regulated during development, such that 99 % of the GluR2 subunit is in the arginine form in the postnatal stage (Dingledine, Borges et al. 1999).

Alternative splicing affects the 38 amino acid sequence in the extracellular S2 domain preceding the C-terminal transmembrane domain M4 of each of the four subunits (Dingledine, Borges et al. 1999). The small segment exists in two alternatively spliced forms referred to as Flip and Flop which are encoded by adjacent exons of the receptor genes. The Flip form is expressed prenatally while the Flop variants are expressed postnatally. These variants can influence the rate and effect of desensitization of heteromeric AMPA receptors and also their sensitivity to specific allosteric modulators (Dingledine, Borges et al. 1999).

AMPA receptors lacking edited GluR2 receptors are expressed in the hippocampus and amygdala. These receptors demonstrate high  $Ca^{2+}$  permeability and predominate in inhibitory interneurons, while non-edited,  $Ca^{2+}$  impermeable GluR2 receptors are found in pyramidal cells (Geiger, Melcher et al. 1995). Extensive evidence in cerebellar stellate cells has shown that activity-induced  $Ca^{2+}$  influx through GluR2-lacking AMPA receptors regulates the targeting of GluR2-containing AMPA receptors, suggesting the existence of a self-regulatory mechanism influencing  $Ca^{2+}$  permeability at the synapses (Liu and Cull-Candy 2002).

#### 2.3 Kainate Receptors

Five subunits of the kainate family of receptors have been cloned: GluR5, GluR6, GluR7, KA1, and KA2 (Bleakman, Gates et al. 2002). These receptors have been subdivided into two families based not only on sequence homology but also agonist-binding properties. GluR5-7 are approximately 70 % identical with each other and 40 % homologous with AMPA receptors (Bleakman, Gates et al. 2002). KA1 and KA2 also share 70 % amino acid sequence homology with each other but only 40 % with GluR5-7. GluR5-7 have a lower affinity kainate-binding site (K<sub>D</sub> of 50–100nM), whereas KA1 and KA2 c have a higher affinity kainate-binding site (K<sub>D</sub> of 5–15nM) (Bleakman, Gates et al. 2002).

Kainate receptors are tetrameric with GluR5-7 forming homomeric and heteromeric combinations. However, KA1 and KA2 cannot form functional homomeric receptors, but combine to form heteromeric assemblies with GluR5-7 (Alt, Weiss et al. 2004). Interestingly, KA2 is not expressed on the cellular surface in the absence of GluR5-7, but is translocated to the endoplasmic reticulum, suggesting the existence of an endoplasmic reticulum signal that is shielded in the presence of other kainate receptors (Gallyas, Ball et al. 2003).

Similar to AMPA receptors, GluR5-7 are susceptible to RNA editing and alternative splicing to introduce further heterogeneity. GluR5-6 but not GluR7 mRNA can be edited at the Q/R site on the M2 segment (Dingledine, Borges et al. 1999). Analogous to AMPA receptors, the addition of arginine can foster lower permeability to  $Ca^{2+}$ . However, unlike GluR2, RNA editing appears to be incomplete during aging and a large number of both edited and non-edited GluR5-6 receptors are distributed in the adult brain (Dingledine, Borges et al. 1999).

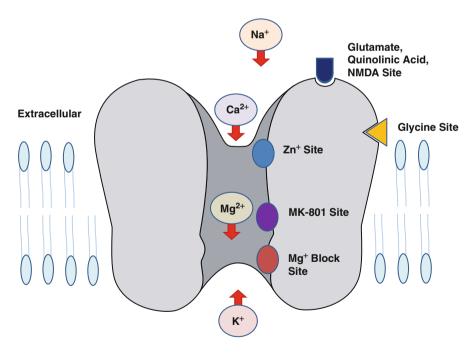
Kainate receptors are distributed throughout the CNS (Bowie and Mayer 1995; Bleakman, Gates et al. 2002). The KA1 and KA2 subunits are both localized in hippocampal CA3 and dentate granule neurons; however, KA2 is more dominant. GluR6 appears to be the major subunit in both presynaptic and postsynaptic receptors at mossy fiber synapses in the hippocampal CA3 region (Kerchner, Wilding et al. 2002). The GluR5 subunit appears to be predominant in the dorsal root ganglion cells (Contractor, Swanson et al. 2000; Kerchner, Wilding et al. 2002). GluR5 is expressed predominantly in hippocampal interneurons where it plays a critical role in regulating the effects of glutamate released by astrocytes (Liu, Xu et al. 2004). GluR5 is also strongly expressed in the temporal lobe. Functional studies have shown that GluR5-containing receptors are involved in synaptic excitation in the amygdala (Rogawski, Gryder et al. 2003).

#### 3 Pharmacology of Ionotropic Glutamate Receptors

Several iGluR agonists and antagonists have been identified. Information regarding the structure and function of these ligands has helped to increase our body of understanding regarding the pharmacology of these receptors and markedly increased our knowledge of receptor physiology and pathophysiology in the CNS.

#### 3.1 NMDA Receptors

NMDA is the most common agonist at the glutamate-binding site (located on the NR2 subunit) of the NMDA receptor (Fig. 4). While it demonstrates lower potency than glutamate, it is not quite a full agonist and shows little selectivity between the NMDA receptor subtypes (Addae, Evans et al. 2000). Another selective full agonist at NMDA receptors is quinolinic acid (QUIN) (De Carvalho, Bochet et al. 1996). QUIN is an endogenous metabolite of the kynurenine pathway (KP). In the CNS, the KP represents the principal route of L-tryptophan metabolism, leading to the formation of the essential pyridine nucleotide nicotinamide adenine dinucleotide (NAD<sup>+</sup>) (Stone 2001). During neuroinflammation, increased activity of the primary KP enzyme indoleamine 2,3-dioxygenase (IDO-1) directs the KP to stimulate



**Fig. 4** The NMDA receptor showing binding sites for both glutamate and glycine. The ligandgated ion channel facilitates movement of calcium, sodium, and potassium across the postsynaptic membrane

QUIN production by activated microglia and invading macrophages, leading to neuronal and glial cell death (Guillemin 2003; Guillemin, Smythe et al. 2004).

Other NMDA receptor agonists include L-aspartate, homocysteate, and homoquinolinate, while *cis*-2,3-piperidinedicarboxylic acid is a low efficacy agonist (Jane, Olverman et al. 1994; Christie, Jane et al. 2000). Additional compounds demonstrate agonist or partial agonist activity at the glycine co-agonist site on the NR1 subunit. Vinyl substituted glycine derivatives such as *S*-hydroxyvinyl glycine can also act as agonists at the glycine site (Kemp, Foster et al. 1988; Priestley, Horne et al. 1990; Priestley, Ochu et al. 1994; Priestley, Laughton et al. 1995). While D-serine is an agonist at low concentrations, it can act as a competitive antagonist at the glutamate site at high concentrations. Other glycine-site partial agonists include L-alanine and D-cycloserine (Kemp and Leeson 1993).

Continuous activation of the NMDA receptor can lead to increased intracellular Ca<sup>2+</sup> permeability and conductance which can trigger a cascade of events leading to either apoptosis or necrosis. The downstream effectors include free radical production, mitochondrial membrane depolarization, and caspase activation (Chen, Harris et al. 1995; Colwell, Altemus et al. 1996; Catts, Ward et al. 1997; Araque, Sanzgiri et al. 1998; Budd, Tenneti et al. 2000; Comoletti, Muzio et al. 2001). Given the importance of NMDA receptors in normal physiological function and in the etiology of neurodegenerative diseases, several NMDA receptor antagonists have been developed to target the glutamate and glycine recognition sites and the NMDA ion channel pore. An example of a high-affinity competitive antagonist of the glutamate site is (R)-2-amino-5-phosphonopentanoate (D-AP5) (Jane, Olverman et al. 1994). While this compound shows high in vivo affinity due to the appropriately placed  $\omega$ phosphonic acid group, it cannot penetrate the blood-brain barrier (BBB) due to its highly charged nature (Kew and Kemp 2005). Recently, adenosine triphosphate (ATP) has been shown to be a dual modulator of the NMDA receptor. While ATP can antagonize the NMDA receptor at the glutamate site at low glutamate concentrations, it can act simultaneously to enhance NMDA function at high glutamate concentrations (Kloda, Clements et al. 2004).

Similarly, numerous inhibitors have been identified at the glycine recognition site (Bergeron, Meyer et al. 1998). Kynurenic acid (KA), a metabolite of the KP, was the first glycine site antagonist described and displayed weak inhibition of the NMDA receptor against QUIN toxicity. In one study, the required concentration of KA to display neuroprotection was tenfold greater than the agonist, QUIN (Jhamandas, Boegman et al. 2000). Many synthetic antagonists of the glycine site were later developed using KA as the pharmacophore (Hartley, Monyer et al. 1990; Foster, Kemp et al. 1992; Kemp and Leeson 1993; Dannhardt and Kohl 1998; Honer, Benke et al. 1998). These compounds include synthetic high-affinity derivatives such as quinolines, quinolones, and indoles. However, compounds of this class can bind tightly to plasma protein which compromises their ability to cross the BBB and produce potent in vivo activity (Bristow, Hutson et al. 1996).

The most potent NMDA antagonists used to treat neurodegenerative diseases are the NMDA ion channel pore blockers. Following the identification of ketamine and phencyclidine as selective NMDA receptor antagonists, the highly potent in vivo active antagonist dizocilpine (MK-801) was developed (Foster, Gill et al. 1987, 1988; Farlow 2004). However, long-term NMDA receptor inhibition by MK-801 has previously been shown to be toxic to cultures of rat cortical neurons (Hwang, Kim et al. 1999). Subsequently, lower affinity compounds such as 1,3-dimethyl-5-aminoadamantane (Memantine) have been developed and approved for clinical use (Farlow 2004). These compounds act as open channel blockers and are irreversibly trapped within the closed pore. Memantine has quicker reversibility, making it efficacious and better tolerated than MK-801 (Kew and Kemp 2005).

A more recent development in NMDA receptor antagonists has been the identification of allosteric modulators that can inhibit the receptor by an interaction with the extracellular N-terminal domain of the NR2 subunit (Kew and Kemp 2005). Compounds such as Ifenprodil and Benzimidazole have high NR2 affinity compared with other NMDA subtypes but produce additional off-target effects by modulating adrenergic receptors and human ether-a-go-go (hERG) ion channels (Finlayson, Witchel et al. 2004). Recently, a phase III clinical trial for Troxoprodil used for the treatment of traumatic brain disease, failed due to poor efficacy (Kew and Kemp 2005).

#### 3.2 AMPA Receptors

AMPA receptors have also been implicated in excitotoxicity due to their ability to increase intracellular  $Ca^{2+}$  overload (Borges and Dingledine 1998). Apart from AMPA itself, several agonists have been developed including ibotenic acid, quisqualic acid, and willardine (Stensbol, Madsen et al. 2002). While the distal acidic group varies between agonists, the amino acid moiety is highly conserved. Unlike glutamate and AMPA which are full agonists and induce rapidly desensitizing responses, kainate is a partial agonist with demonstrated low desensitization. The degree of desensitization has been linked to the extent of dimerization between pairs of subunits within the AMPA receptor (Stensbol, Madsen et al. 2002).

The first competitive AMPA receptor antagonists were 6-cyano-7-nitroquinoxaline-2,3-dione (CQNX) and 6,7-dinitro-quinoxaline-2,3-dione (DNQX). However, these compound showed low selectivity and could also antagonize the glycine-binding site of the NMDA receptor. More selective AMPA receptor antagonists include 2,3-dihydroxy-6-nitro-7sulfamoyl-benzo(F)quinoxaline (NBQX) (Birch, Grossman et al. 1988; Turski, Huth et al. 1998). However, these compounds show poor water solubility and represent failed clinical trials due to adverse side effects including nephrotoxicity. Several noncompetitive AMPA receptor antagonists have been developed. These include several 2,3-benzodiazepine compounds and quinazoline derivatives. These compounds are selective for AMPA over NMDA and kainate receptors and have recently entered clinical trials (Lazzaro, Paternain et al. 2002; Solyom and Tarnawa 2002).

Additionally, several AMPA-receptor-positive allosteric modulators have been developed. These new class of compounds can modulate receptor desensitization/deactivation without activating the receptor when supplied alone (Danysz 2002). These compounds include cognitive enhancing "nootropic agents" such as 1-(4-methoxybenzoyl)-2-pyrrolidinone (aniracetam), benzamides, and benzothiadiazides. These compounds demonstrate distinct mechanistic profiles and are selective for individual AMPA receptor subunits (Danysz 2002).

## 3.3 Kainate Receptors

Kainate receptors have also been associated with increased  $Ca^{2+}$  influx, leading to excitotoxic neuronal cell death. Apart from kainate, several kainate-receptor agonists have been identified. These include 5-tert-butyl-4-isoxazolepropionic acid (ATPA), a substituted analogue of AMPA, and (*S*)-5-iodowillardine (Lerma, Paternain et al. 2001; Bleakman, Gates et al. 2002). These compounds show lower affinity to AMPA receptors. The quinoxalinedione compounds such as CNQX and NBQX are competitive antagonists at kainate receptors but show little discrimination between kainate and AMPA receptors (Lerma, Paternain et al. 2001; Bleakman, Gates et al. 2002). Derivatives of willardine also exhibit antagonist effects at both AMPA and kainate receptors. Kainate receptors also demonstrate rapid desensitization following agonist binding (Bleakman, Gates et al. 2002; Alt, Weiss et al. 2004). The lectin concanavalin A can rapidly inhibit receptor desensitization by interacting with *N*-glycosylated residues and increase receptor agonist affinity. It also exhibits activity at NMDA and AMPA receptors (Lerma, Paternain et al. 2001; Bleakman, Gates et al. 2002).

# 4 Molecular Mechanisms in Neurodegenerative Diseases

There is an attractive body of evidence incriminating glutaminergic dysfunction in the process of slow excitotoxicity associated with neuronal loss in Alzheimer's disease (AD), Parkinson's disease (PD), Huntington's disease (HD), and Amyotrophic Lateral Sclerosis (ALS).

# 4.1 Alzheimer's Disease

Alzheimer's disease (AD) is a progressive neurodegenerative disorder associated with the progressive loss of cholinergic neurons parallel to impairments in cognition and memory (Mattson 2004). The pathological hallmarks of AD include extracellular deposition of the  $\beta$ -amyloid peptide (A $\beta$ ) in senile plaques and formation of intracellular neurofibrillary tangles (NFTs) (Hartmann 1999). Accordingly, the A $\beta$  hypothesis suggests that AD occurs as a response to increased production and extracellular deposition of A $\beta$  and intracellular deposition of NFTs. These abnormal pathologies can trigger numerous neurotoxic deleterious processes including excitotoxicity, oxidative stress and energy depletion, neuroinflammation, and cell death (Bamberger and Landreth 2001). Interestingly, the glutaminergic system is severely compromised by A $\beta$ -mediated regulation of glutamate receptors in various brain regions, correlating with significant cognitive impairment (Parameshwaran, Dhanasekaran et al. 2008).

The neuronal degeneration in AD has been linked to overactivation of the NMDA receptor (Butterfield 1997). Recent studies have shown that A $\beta$  can stimulate free radical formation through a mechanism involving NMDA receptors. QUIN can increase mRNA and protein expression of the inducible (iNOS) and neuronal (nNOS) forms of nitric oxide following overactivation of the NMDA receptor, leading to subsequent induction of nitric oxide synthase in human primary fetal astrocytes and neurons, triggering cell death (Braidy, Grant et al. 2009). Dysregulation of NMDA receptor activation and free radical formation may serve as dual deleterious roles in AD.

We have previously shown that QUIN is significantly increased in postmortem AD brain (Guillemin, Brew et al. 2007). Recently, our group has shown that the KP is activated in the AD brain with QUIN immunoreactivity reported in cortical microglia, astrocytes, and neurons, and in the perimeter of A $\beta$  plaques and NFTs. We recently reported an accumulation of QUIN in dystrophic neurons in cortical sections from AD patients (Guillemin, Brew et al. 2005). Additionally, QUIN can increase tau phosphorylation and stimulates dysregulation of microtubule formation in vitro. We also found that QUIN can induce at least ten important neuronal genes strongly associated with AD pathology (Rahman, Ting et al. 2009). Therefore, excitotoxicity due to increased activation of NMDA receptors can enhance the vulnerability of local neurons consistent with AD pathology.

Restoration of homeostasis in the glutamatergic system using NMDA receptor antagonists has been shown to ameliorate the neurotoxic effects by NMDA activation both in vitro and in vivo. Memantine, a noncompetitive NMDA receptor channel blocker, has received marketing approval from the EMEA (European Medicines Agency), FDA (Food and Drug Administration), and TGA (Therapeutic Goods Administration) for the treatment of mild-to-severe AD in Europe, USA, and Australia (Doggrell 2003). It has strong voltage-dependency and fast kinetics to help provide neuroprotection and reverse deficits in memory and learning (Ferris 2003; Gallarda and Loo 2004; Kato 2004; Gardoni and Di Luca 2006).

#### 4.2 Parkinson's Disease

Parkinson's disease (PD) is a neurological disease caused by degeneration of dopaminergic neurons in the substantia nigra, leading to a significant reduction in dopamine levels in the striatum (Mosley, Benner et al. 2006). Although the exact etiology of PD remains unknown, excessive glutaminergic drive has been proposed as a major source of excitotoxicity in the nigra (Barnum and Tansey 2010). Studies have shown that parkin (hereditary Parkinson disease PARK2 gene product) can modulate the stability and function of excitatory glutaminergic synapses (Yu, Sun et al. 2011). Postsynaptic expression of parkin can inhibit excitatory synaptic

transmission. Knockdown of parkin or PD-linked parkin mutants can promote synaptic function and facilitate the proliferation of glutamatergic synapses. Activation of the NMDA receptor can lead to increased  $Ca^{2+}$  influx which has been linked to cell death in PD (Yu, Sun et al. 2011). This is further supported by the observation that dopaminergic neurons expressing the calcium-binding protein calmodulin are selectively preserved in the PD brain (Pierson, Norris et al. 2004).

Several studies have shown that excitotoxicity plays a key role in toxicity induced by 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) toward dopaminergic neurons (Araki, Kumagai et al. 2000). MPTP is converted to the 1-methyl-4-phenylpyridinium ion (MPP<sup>+</sup>) in glial cells by monoamine oxidase-B. The active metabolite is taken up by the dopamine transporter into dopaminergic neurons where it has been shown to inhibit mitochondrial complex I (Beal 2003). This not only compromises neuronal energy production, but also induces a slow excitotoxicity, leading to the loss of nigral cells. Inhibition of the glutamatergic corticostriatal tract has been shown to reduce the size of MPP+-mediated lesions in the rat striatum. Furthermore, the glutamate-release inhibitor riluzole has been shown to inhibit glutamate release in the corticostriatal tract to attenuate intrastriatal administration of MPP<sup>+</sup> on dopaminergic neurons (Araki, Kumagai et al. 2000). MK-801 (dizocilipine), a noncompetitive NMDA receptor antagonist, has been shown to protect against the toxicity of MPTP in a rat model for PD (Hsieh, Gu et al. 2012). NBQX, an AMPA/kainate-receptor antagonist, has been shown to prevent the symptoms of PD, but not the neurodegeneration in MPTP-treated monkeys (Luqquin, Obeso et al. 1993). The finding that MK-801 cannot protect against MPP<sup>+</sup> toxicity in primary cultures of dissociated mesencephalic neurons suggests that alternative mechanisms of action for MPTP may exist in vivo.

#### 4.3 Huntington's Disease

Huntington's disease (HD) is an inherited neurodegenerative disorder characterized by cognitive and psychiatric impairment, and dyskinesia (Beal, Kowall et al. 1986). One of the major hypotheses developed to explain the vulnerability of medium-size spiny neurons (MSNs) in the neostriatum is the excitotoxicity hypothesis. Studies performed as early as the 1970s in rats, mice and other rodents demonstrated that striatal kainate injections can destroy MSNs while sparing neighboring axons and the effect is similar to HD pathology observed in humans (Ferrante, Kowall et al. 1993; Figueredo-Cardenas, Anderson et al. 1994; Estrada Sanchez, Mejia-Toiber et al. 2008). Similarly, striatal injections of QUIN in rats are able to reproduce the lesions reported in HD (Ellison, Beal et al. 1987; Bruyn and Stoof 1990; Ceresoli-Borroni, Guidetti et al. 1999). QUIN-induced lesions in experimental rodent models can be attenuated using NMDA receptor antagonists (Ellison, Beal et al. 1987; Bruyn and Stoof 1990; Ceresoli-Borroni, Guidetti et al. 1990). The QUIN model has been extended to primates where similar pathological hallmarks have been observed together with clinical symptoms of chorea (Bruyn and Stoof 1990).

Intrastriatal infusion of kainate has also been shown to induce excitotoxic lesions in the striatum and induces neuronal apoptosis via activation of NF- $\kappa$ B and induction of p53 and other pro-apoptotic proteins (Nakai, Qin et al. 2000). However, AMPA/ kainate-receptor antagonists have been shown to be inactive in these models (Levine, Klapstein et al. 1999).

The HD gene and its protein product huntingtin (htt) have been proposed to play a key role in neuronal cell death. It has been demonstrated that mutant expression of htt can stimulate NMDA receptor activity to disrupt intracellular calcium homeostasis (Zhang, Li et al. 2008). Recent studies have shown that the postsynaptic density protein PSD-95 represents a link between htt and NMDA receptor function. One study has shown that PSD-95 can bind to the C-terminal of the NR2 subunit to stabilize the NMDA receptor (Roche, Standley et al. 2001). Co-immunoprecipitation studies have also revealed interactions between PSD-95 and NMDA receptor subunits NR1, NR2A, and NR2B, thus providing further evidence for the association of NMDA receptors with PSD-95 (Sun, Savanenin et al. 2001).

#### 4.4 Amyotrophic Lateral Sclerosis

Amyotrophic Lateral Sclerosis (ALS) is characterized by progressive muscle weakness, leading to paralysis and death, generally due to respiratory failure. Histologically, ALS is characterized by the selective loss of both upper (corticospinal) and lower (spinomuscular) motor neurons (Boillee, Vande Velde et al. 2006; Boillee, Yamanaka et al. 2006). There are several lines of evidence to support the role of excitotoxicity in the loss of motor neurons. Intrathecal injections of kainate have been shown to induce neuronal damage in the spinal cord (Hugon, Ludolph et al. 1988; Hugon and Vallet 1990). These lesions are characterized by axonal swelling and impaired cytoskeletal architecture similar to those seen in ALS (Hugon and Vallet 1990). In chicken motor neurons, glutamate-initiated neurotoxicity is predominantly due to AMPA/kainate receptors. However, in rats, excitotoxicity is mediated by both NMDA and AMPA/kainate receptors (Estevez, Stutzmann et al. 1995; Carriedo, Yin et al. 1996). The observed toxicity in these cell types can be attenuated using AMPA/kainate-receptor antagonists but not NMDA receptor antagonists (Rothstein 1995). However, in humans, we have shown that OUIN can induce apoptosis of fetal motor neurons in a dose-dependent manner and the effect is ameliorated by blocking the NMDA receptor (Chen, Brew et al. 2011).

If excitotoxicity is indeed a major pathogenic mechanism in ALS, it is necessary to explain why only certain motor neurons are affected. A major factor determining the vulnerability of certain motor neurons to excitotoxicity is associated with the subunit composition of the iGluR that they may express. In situ hybridization studies have shown that the NR2B NMDA receptor subunit is absent in rat motor neurons while the GluR2 AMPA/kainate receptor is not present in human spinal motor neurons (Tolle, Berthele et al. 1993). In the absence of the GluR2 subunit, AMPA/kainate receptors are more permeable to  $Ca^{2+}$  and this may explain the excitotoxicity observed in rats (Williams, Day et al. 1997). Nonetheless, direct electrophysiological studies investigating intracellular calcium influx into motor neurons resulting from AMPA/kainate-receptor-mediated excitotoxicity have not been investigated and no comparison has been made between iGluR in rats and humans.

Although no effective cures have been developed, anti-excitotoxic drugs have provided the first treatment for this debilitating disorder. Early trials conducted using the noncompetitive NMDA receptor antagonist dextromethorphan have shown negative results (Hollander, Pradas et al. 1992). Given that dextromethorphan is a weak NMDA receptor antagonist at the doses used (<150 mg/day), it may be ineffective in achieving complete receptor blockade (Beaton, Stemsrud et al. 1992). Moreover, pharmacokinetic studies have shown that side effects of the drug occur at doses five times higher than those used in the efficacy studies (Hollander, Pradas et al. 1994). Plasma levels of dextromethorphan necessary for significant NMDA receptor blockade were unlikely to be reached in the clinical trials.

Promising results have been obtained with the glutamate-release inhibitor and sodium channel blocker, riluzole. A modest increase in survival was observed following 1 year of treatment which was associated with a significant reduction in the rate of deterioration in motor function (Zoccolella, Beghi et al. 2007). Another trial confirmed the beneficial effects of riluzole on survival but not on motor function (Gurney, Fleck et al. 1998). Accordingly, riluzole has been approved and marketed for the treatment of ALS in various countries (Miller, Mitchell et al. 2007). While this is encouraging to those who argue that blockade of excitotoxicity is an important therapeutic strategy to treat neurodegenerative diseases, it remains to be demonstrated whether riluzole is neuroprotective in ALS.

#### 5 Conclusion

The concept of excitotoxicity is implicated in the pathogenesis of a variety of neurodegenerative diseases. Neurodegenerative diseases may develop as a result of abnormally high concentrations of excitotoxic amino acids, or by a more insidious process whereby energetically weakened neurons become vulnerable to ambient concentrations of glutamate or an exogenous congener. Currently available anti-excitotoxic drugs have played vital roles in examining the factors which facilitate neuronal death in several experimental paradigms. Where suitable animal models are available, such as for AD and PD, iGluR inhibitors have paved the way for defining the etiology and treatment of neurodegenerative diseases in the clinic. The next decade will lead to the development of newer and more specific compounds to block excitatory neurotransmission, and may include NMDA and AMPA/kainate-receptor antagonists. The current century may witness a revolution in the treatment of neurological diseases using anti-excitotoxic drugs.

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# **Glutamate Neurotoxicity, Transport and Alternate Splicing of Transporters**

# Aven Lee, Shannon Beasley, and David V. Pow

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#### Abstract

Glutamate is the major excitatory neurotransmitter in the central nervous system and its levels in the synaptic cleft are tightly controlled by high affinity glutamate transporters (also known as *Excitatory Amino Acid Transporters* or EAATs). The EAAT family is comprised of five members (EAAT1-5), and these transporters are subject to alternative splicing. Alternative splicing of the EAAT genes is a fundamental mechanism that can give rise to multiple distinct

A. Lee (🖂) • S. Beasley

D.V. Pow

School of Medical Sciences and Health Innovations Research Institute, RMIT University, Melbourne, VIC, Australia

e-mail: david.pow@rmit.edu.au; david@powfamily.com

UQ Centre for Clinical Research, The University of Queensland, Brisbane, QLD, Australia e-mail: aven.lee@uq.edu.au

mRNA transcripts, producing protein isoforms with potentially altered functions. Numerous splice variants of EAATs have been identified in humans, rodents, and other mammalian species. All splice variants of EAATs cloned to date are either exon-skipping and/or intron-retaining types. These modifications may impact upon protein structure, posttranslational modification, function, cellular localization, and trafficking. Message and protein for these splice variants are detectable in the normal brain and, in many instances, have been shown to be induced by pathophysiological insults such as hypoxia. In addition, aberrant expression of EAAT splice variants has been reported in neurodegenerative conditions such as amyotrophic lateral sclerosis, Alzheimer's disease, ischemic stroke, and agerelated macular degeneration. These EAAT variants may represent therapeutic targets and thus require an improved understanding of their regulation. This chapter describes recent developments in investigating the molecular heterogeneity, localization, function, structure, and regulation of the EAATs and their splice variants.

#### Keywords

EAAT • GLAST • GLT1 • EAAC1 • Excitotoxicity • Splicing • Stroke • Schizophrenia • Motor neurone • ALS • Glaucoma

List of Abbreviations					
AD	Alzheimer's disease				
ALS	Amyotrophic lateral sclerosis				
AMD	Age-related macular degeneration				
CNS	Central nervous system				
EAAC1	Excitatory amino acid carrier 1				
EAAT	Excitatory amino acid transporter				
GABA	Gamma-aminobutyric acid				
GLAST	Glutamate aspartate transporter				
GLT-1	Glutamate transporter-1				
GTRAP	Glutamate transporter associated protein				
hnRNP	heterogeneous nuclear ribonucleoproteins				
HSP70	70 kDa heat shock proteins				
MAP2	Microtubule-associated protein 2				
MLO-Y4	Murine long bone osteocyte Y4				
mRNA	messenger ribonucleic acid				
PDZ	Postsynaptic density (PSD), Drosophila disc large tumor				
	suppressor (Dlg1), and zonula occludens-1 protein (ZO-1)				
SLC	Solute carrier				
SOD	Superoxide dismutase				
SR protein	Serine/arginine-rich protein				
SRPK	Serine/arginine-rich protein kinase				
UTR	Untranslated region				

#### 1 Introduction

#### 1.1 Neurotransmission: A Regulated Process of Neurotransmitter Release and Removal

The brain is a highly regulated environment, protected by a blood-brain barrier and other homeostatic mechanisms, to ensure that events such as neurotransmission are able to occur without impediment. Neurotransmission is typically mediated by the evoked fusion of one or more synaptic vesicles containing around 10,000 molecules of neurotransmitter, each of which can potentially activate a receptor molecule. Thus, the timeframe for the neurotransmission signaling event is heavily dependent upon the kinetics of clearance of the neurotransmitter. Accordingly, an understanding of the biological events that influence the kinetics of neurotransmitter removal from the extracellular space is central to our understanding of brain function.

Many neurotransmitters are removed from the extracellular space using high affinity neurotransmitter transporters, and these are frequently pharmacological targets. Thus, the serotonin transporter is a target for selective serotonin reuptake inhibitors (SSRIs) used to treat disorders such as depression. This whole area of pharmacology depends upon the drug interacting with a defined transporter that is expressed in a predictable manner with predictable properties. However, alternately spliced forms of many high affinity neurotransmitter transporters have been identified, including the serotonin transporter (Sakai et al. 2003), dopamine transporter (Talkowski et al. 2010), noradrenalin transporter (Sogawa et al. 2007), betaine-GABA transporter (Takenaka et al. 1995), and glycine transporters (Ponce et al. 1998; Hanley et al. 2000). In the case of the serotonin transporter, there is evidence for differential transport capacity of serotonin by the variant forms, while alternately spliced dopamine transporter may influence relative expression of the functional form of the transporter and thus perturb extracellular levels of dopamine. These types of finding lead to the conclusion that alternate splicing might be a mechanism that differentially regulates clearance of many neurotransmitters under differing physiological and pathophysiological states.

This chapter focuses upon the most abundant excitatory neurotransmitter in the central nervous system (CNS), namely, glutamate because glutamate is a molecule that may be dysregulated in a wide variety of CNS disease states, ranging from Alzheimer's disease, schizophrenia, motor neuron disease, and glaucoma to depression (Cowburn et al. 1988; McCarthy et al. 2012; Moghaddam and Javitt 2011). Therefore, an understanding of the transporters that mediate glutamate removal and the potential ways in which the transporters might be modified by events, such as alternate splicing, are central to our understanding of brain pathophysiology and ultimately to the development of therapeutic paradigms that optimize brain glutamate homeostasis. Extracellular and intracellular levels of

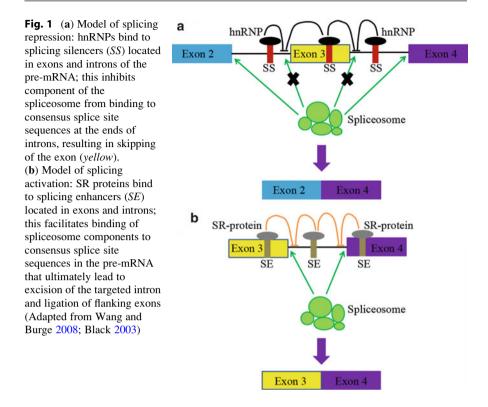
glutamate are regulated principally a family of high affinity sodium-dependent Excitatory Amino Acid Transporters (EAATs), which are members of the Solute Carrier Family-1 (SLC1) (Danbolt 2001). In the mammalian CNS, five major subtypes of EAATs have been identified; they share considerable homology (50-60 %) at the amino acid level, but each member exhibits distinct and differential localizations at the subcellular, cellular, and regional levels. EAAT1 (SLC1A3), the human homologue of the rodent glutamate/aspartate transporter (GLAST), is present in glia throughout the CNS but is particularly abundant in the radial (Bergmann) glia of the cerebellum and the radial (Müller) glia of the retina; EAAT2 (SLC1A2), the human homologue of glutamate transporter 1 (GLT-1), is present mainly in glia, particularly astrocytes, and is widespread throughout the CNS; EAAT3 (SLC1A1), the human homologue of Excitatory Amino Acid Carrier 1 (EAAC1), is localized to subsets of neurons throughout the CNS; EAAT4 (SLC1A6) is primarily localized to cerebellar Purkinje cells but with some expression in astrocytes (Hu et al. 2003) and EAAT5 (SLC1A7) is abundant in the retina, particularly in photoreceptors and populations of bipolar cells (Reviewed by Beart and O'Shea 2007).

One mechanism that is likely to influence EAAT function or dysfunction is alternative splicing. Alternative splicing of RNA is a common mechanism that allows a single gene to produce multiple different protein products (Lopez 1998). This versatile mechanism can also regulate parameters such as the amount of mRNA expressed and the stability of such mRNA. Changes in encoded protein sequence can affect nearly all aspects of protein function, including protein stability, binding properties, posttranslational modifications, cellular localization, and trafficking (Stamm et al. 2005). In recent years, many splice variants of EAATs have been described in humans, rodents, and other mammalian species. Message and protein for these splice variants are detectable in the normal brain and in some instances, their expression can be induced by pathophysiological insults. This chapter reviews recent progress in investigating the molecular heterogeneity, localization, function, structure, and regulation of the EAATs and their splice variants.

# 2 Splicing of Glutamate Transporters

#### 2.1 Mechanism of Splicing

Alternative splicing of EAATs arises from differential removal of exons from the pre-mRNA that was transcribed from the DNA, or in some instances the retention of introns from the pre-mRNA, to yield many distinct mRNA and protein isoforms. Exon skipping is a common mode of splicing in mammalian brain, generating, for instance, the four known forms of GLAST (EAAT1). By contrast, GLT-1 (EAAT2) is spliced in a combinatorial manner (exon skipping and/or intron retention) to produce many distinct protein isoforms (for review, see Lee and Pow 2010).



It is notable that alternate splice variants for EAAT3 and EAAT4 have not so far been described in the mammalian brain, though single nucleotide polymorphisms are undoubtedly present. However, given the limited attention that has been paid to these transporters to date, it is plausible that splice variants may exist but have not yet been detected, and this remains to be resolved. A similar situation has recently been resolved for EAAT5, which had until recently been described in mammals as a single form of the protein. Lee et al. (2012a) have recently demonstrated that it also exists as multiple splice variants, and that all splice variants cloned to date are either exon-skipping or intron-retaining types.

The regulation of EAAT splicing involves consensus splice site sequences in the pre-mRNA and cellular factors such as the "spliceosomes" which are dynamic complexes of proteins and RNA that act as ribozymes, to typically excise introns and ligate the flanking exons. In addition to the consensus splice site sequence on the pre-mRNA, auxiliary elements such as splicing enhancers and splicing silencers can strongly influence alternative splicing; such elements are variable in sequence and contain binding sites for serine/arginine-rich (SR)-proteins or heterogeneous nuclear ribonucleoproteins (hnRNPs), the binding of which can activate or repress the adjacent splice sites, respectively (Fig. 1). Therefore, the binding of

SR proteins and hnRNPs to such elements strongly influences the choice of splicing sites and regulate the formation of active spliceosomes in the pre-mRNA. Whether an exon is included in the mature transcript is determined by the concentration or activity of each type of regulators (SR proteins and hnRNPs). Concentration of SR proteins and hnRNPs can vary significantly between different cell types, and it is thought that such differential expression of SR proteins and hnRNPs affects the splicing of many pre-mRNAs (for reviews, see Chen and Manley 2009; Matlin et al. 2005).

The activity of SR proteins is strongly influenced by the phosphorylation state of these proteins. At least four different protein kinase families have been shown to phosphorylate SR proteins. The most specific of these is the SRPK family (e.g., SRPK1 and SRPK2), which are typically localized to the cytoplasm of cells, indicating that regulation of SR proteins is a cytoplasmic event. SR protein kinases are also regulated; SRPK1 binds to co-chaperones including heat shock proteins such as HSP40, in turn influencing interactions with HSP70 and HSP90. Accordingly, stress responses in cells such as astrocytes and neurons, including production of heat shock proteins, may modify pre-mRNA splicing for transporters such as GLT-1 (EAAT2). The activity of hnRNP proteins can also be modified by phosphorylation. For example, hnRNP A1 is directly phosphorylated by PKA, PKC, and Casein Kinase II (reviewed in Lynch 2007). Serine/threonine phosphorylation under hypoxia and stress conditions (van der Houven van Oordt et al. 2000; Daoud et al. 2002; Allemand et al. 2005) was previously shown to cause hyperphosphorylation and accumulation of hnRNPs in the cytosol. This could serve the purpose of removing such regulatory factors from pre-mRNA processing events. Indeed, a number of in vivo studies have clearly shown that splice site selection is altered following hypoxia and application of osmotic shock (van der Houven van Oordt et al. 2000; Daoud et al. 2002). Accordingly, stressors such as hypoxia are likely to affect the nuclear ratio of specific splicing regulators (SR proteins and hnRNPs) in cells such as astrocytes, thereby varying the regulation of alternative splicing of particular EAATs.

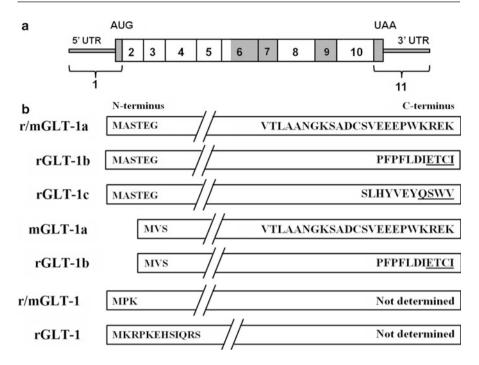
# 2.2 Splicing of 5'-Untranslated Region

Alternative splicing is a mechanism used for generating 5'-heterogeneity of transporters such as GLT-1 (EAAT2). Multiple alternative 5'-untranslated regions (UTR) exist for GLT-1 in human, rat, and mouse, with studies showing that mRNA levels for such variants change following pathological insults. Tian et al. (2007) showed that GLT-1 variants with longer 5'-UTRs (>310 nucleotides) were highly regulated at the translational level by factors such as corticosterone and retinol. In addition, some disease-associated insults, such as elevated glutamate levels and oxidative stress, were shown to affect the translational efficiency of such transcripts in primary astrocyte cultures. These observations suggest that expressions of such GLT-1 variants at the mRNA level are highly regulated under normal and disease state conditions and will thus influence GLT-1 protein expression.

Splicing of the GLT-1 gene can produce transcripts with long 3'-UTR's; both human EAAT2 and mouse GLT-1a are known to contain unusually long 3'-UTRs (>9 Kb), suggesting that they contain potential elements that control mRNA localization, translational efficiency, or stability of mRNA (Kim et al. 2003; Kirschner et al. 1994). The 3'-UTR of human EAAT2 is reported to contain 23AU-rich elements (AUUUA) and 8 putative polyadenylation signals that could potentially be used to generate shorter EAAT2 transcripts (Kim et al. 2003); thus, Munch et al. (2000) have reported human EAAT2 transcripts of varying sizes (1.8, 2.4, 7, and 12 Kb) in human brain. Furthermore, sequences at the extreme end of the EAAT2/GLT-1a 3'-UTR is highly conserved between human, rat, mouse, and macaque, suggesting that this region has a regulatory function in posttranscriptional control (Kim et al. 2003). Computational analyses of the 3'-UTR of GLT-1a have also revealed the presence of putative K boxes, which are binding sites for microRNAs such as miR-218 (Lewis et al. 2005). Interestingly, this binding site is absent in the 3'-UTR of another alternately spliced form of GLT-1 called GLT-1b, suggesting that miR-218 may be a specific regulator of GLT-1a (for a detailed review, see Lauriat and McInnes 2007). Alignment analysis of 3'-UTRs between GLT-1a and GLT-1b revealed that there are sequences unique to each variant. These unique regions may contain potential regulatory elements that control the differential expression of the two GLT-1 variants. The complete sequence of the 3'-UTR of a third form called GLT-1c is unknown, and thus, it is not known if this variant contains any unique 3'-UTR sequences. However, it is interesting to note that expression levels of both GLT-1b and GLT-1c are much lower than that of GLT-1a in brain tissue homogenates, raising the possibility that in brain at least, their expression may be more tightly controlled by trans-acting factors (such as RNA-binding proteins) (Lauriat and McInnes 2007). Conversely, the highly abundant expression of GLT-1b and GLT-1c in the retina albeit in distinct populations of neurons, relative to GLT-1a (Pow et al. 2000; Rauen et al. 2004), suggests both a cell-specific enhancement of GLT-1b and GLT-1c expression and a co-associated suppression of expression of GLT-1a in this tissue.

## 2.4 Alternative Splicing and Generation of PDZ-Binding Motifs

Three distinct C-terminal regions of GLT-1 are generated from alternative splicing, and these variants are referred to as GLT-1a, GLT-1b, and GLT-1c. GLT-1a is the wild-type form of GLT-1 cloned by Pines et al. (1992) and contains 11 exons. GLT-1b lacks exon 11 and the coding region terminates in the intron after exon 10, generating a novel 11 amino acid sequence at the C-terminus (Chen et al. 2002; Schmitt et al. 2002). Similarly, GLT-1c includes exons 1 to 10 and terminates with a small exon spliced in from partial sequences of intron 10, again generating a novel 11 amino acid sequence at the C-terminus (Rauen et al. 2004).



**Fig. 2** (a) Exon structure of GLT-1 mRNA with exons subject to alternate splicing depicted in gray. Exons are depicted by *rectangles*; *UTR* untranslated region, *AUG* start codon, *UAA* stop codon. (b) Schematic illustration depicting "normally" spliced GLT-1 (referred to as GLT-1a), and known alternate N-terminal and C-terminal amino acid sequences, with PDZ motifs underlined. "m" prefix denotes sequences identified in mouse; "r" indicates sequences from rat

The last four amino acids in the C-termini of both GLT-1b (ETCI) and GLT-1c (QSWV) conform to the consensus sequence of a class I PDZ-binding motif (X-S/T-X- $\Phi_{COOH}$ ), where X is any residue and  $\Phi$  is a hydrophobic residue (Ivarsson 2012) (Fig. 2). These motifs are similar to the C-terminal PDZ motif (ETKM) present on GLAST that bind to the NHERF1 PDZ protein, which in turn interacts with ERM proteins and the astrocyte cytoskeletal protein GFAP (Lee et al. 2007; Sullivan et al. 2007b). Disruptions of these interactions were shown to perturb glutamate transport (Sullivan et al. 2007b). The PDZ proteins are particularly interesting, because they play key roles in anchoring transmembrane proteins to the cytoskeleton and hold together signaling complexes (Reviewed by Romero et al. 2011). Accordingly, the GLAST-NHERF1 interactions may limit the targeting of this transporter to central "core" zone of astrocytes where GFAP predominates (Sullivan et al. 2007b).

GLT-1b does not interact with NHERF1 (Lee et al. 2007) and potential PDZ-binding partners that associate with this transporter variant in astrocytes are still unclear. Bassan et al. (2008) have shown that GLT-1b can interact with the PDZ protein PICK1; however, very little PICK1 is expressed by astrocytes.

Similarly, Gonzalez-Gonzalez et al. (2008) demonstrated GLT-1b interactions with PSD95; however, this PDZ protein is mainly expressed in neurons, and anatomical co-localization of GLT-1b and PSD95 has not been demonstrated to indicate a functional relationship between the two proteins in brain.

No PDZ-binding partner has been identified for GLT-1c, which is abundantly expressed in photoreceptors of the retina. GLT-1c is also expressed in the brain at low levels, being primarily restricted to end-feet of astrocytes (Rauen et al. 2004). The differential expression of GLT-1c relative to GLT-1b may be indicative of interactions with different PDZ proteins and participation in different signaling pathways.

PDZ-binding motifs are also present on other EAATs, including EAAT3, EAAT4, and EAAT5. Neuronal EAAT4 was previously shown to interact with GTRAP48 (glutamate transporter-4 associated protein), a PDZ-domain-containing protein expressed predominantly in the cerebellum (Jackson et al. 2001). GTRAP48 specifically interacts with the C-terminal PDZ motif (ESAM) of EAAT4 and positively modulates its glutamate transport activity by enhancing the cell surface expression of EAAT4 (Jackson et al. 2001). Similarly, the cell surface density of EAAT3 is regulated via interaction of its C-terminal PDZ motif (TSQF) with the PDZK1 adaptor protein (D'Amico et al. 2010). The retinal EAAT5 has also been reported to contain a PDZ-binding motif (ETNV) at its C-terminus, based on findings from yeast two-hybrid screening, making it likely that the EAAT5 C-terminus may be involved in EAAT5-PDZ interactions (Arriza et al. 1997). The inclusion of PDZ-binding motifs in many of the EAATs suggests that interactions with PDZ-domain-containing proteins are common and probably important for creating differential targeting with consequential functional diversity. It is therefore plausible that splice variants differing in their C-terminal PDZ motif sequences may play distinct roles in regulation of neurotransmission or development.

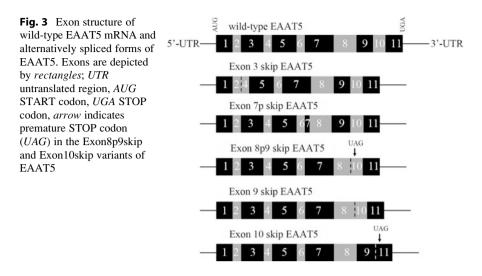
# 2.5 Alternative Splicing and Generation of Exon-Skipping EAAT Variants

Several exon-skipping GLAST (EAAT1) variants are generated from alternative splicing. Huggett et al (2000) identified a splice variant of GLAST, lacking exon 3 (GLAST-1a), which is expressed in brain and bone. MLO-Y4 cells transfected with GFP-tagged GLAST and GLAST-1a appear to target the proteins to different cellular compartments; GLAST was targeted to the plasma membrane whereas GLAST-1a appeared to be expressed within internal vesicles (Huggett et al. 2002). When an antibody was later developed against GLAST-1a, expression of this protein was readily demonstrable in populations of astrocytes throughout the brain and retina (Macnab et al. 2006). There are notable differences between the distribution of GLAST-1a and wild-type GLAST in retina; GLAST-1a is preferentially localized to the outer end-feet of Muller cells, whereas GLAST is distributed throughout the cells, suggesting targeting and compartmentalization to different

parts of the cells. GLAST-1b is an exon 9–skipping form of GLAST and based on preliminary studies in transfected HEK293 cells was suggested to be a nonfunctional glutamate transporter which exerts a dominant negative effect on wild-type GLAST function (Vallejo-Illarramendi et al. 2005). Expression of GLAST-1b is highly upregulated at the protein level (albeit mainly in neurons), following insults such as hypoxia (Sullivan et al. 2007a). GLAST-1c is a splice variant of GLAST, lacking both exon 5 and exon 6 that was cloned from neonatal hypoxic pig brain (Lee et al. 2012b). GLAST-1c protein was shown to be abundantly expressed in astrocytes and oligodendrocytes in rat brain, pig brain, and human brain, including gray and white matter. Similarly, expression of GLAST-1c was observed in primary astrocyte cultures and in cultured oligodendrocytes. In astrocytes, GLAST-1c rapidly redistributed to the cell surface upon stimulation of protein kinase with phorbol esters, suggesting that phosphorylation plays a key role in the targeting of this protein.

Numerous exon-skipping variants of GLT-1 have been identified in human and rodent brains. These include a GLT-1 splice variant, missing exon 7, and a form, missing both exon 7 and exon 9 (Scott et al. 2011). Meyer et al. (1998, 1999) have discovered a form of GLT-1 missing exon 9 as well as more complex splice variants such as a form missing parts of exon 6 and exon 7 and a form that skips between part of exon 7 and part of exon 11. Little information is available on the expression of these proteins. Macnab and Pow (2007a) have generated antibodies against the splice site of the exon 9-skipping form of GLT-1 and demonstrated its expression in populations of white matter astrocytes. More recently, Lee et al. (2011) cloned an exon 4-skipping form of GLT-1 from the rodent brain, and an antibody that selectively recognized this variant revealed strong signals in gray matter astrocytes. Interestingly, the 5'-UTR of the exon 4-skipping GLT-1 is 306-nucleotides longer than that of regular GLT-1 (Pines et al. 1992). This is of interest because Tian et al. (2007) have shown that some GLT-1 transcripts bearing longer 5'-UTR are translationally regulated by factors such as corticosteroids and retinol as well as by several disease-associated insults in primary astrocyte cultures and in mice. Therefore, the possibility exists that translational regulation may play an important role in expression of exon 4-skipping GLT-1 under normal and disease conditions.

EAAT5 also exists as alternatively spliced forms in the retina. By PCR cloning and sequencing, Lee et al. (2012a) recently demonstrated the presence of five novel splice variant forms of EAAT5 which skip either partial or complete exons in the rat retina: Three of the splice variants skipped full exons, resulting in forms of EAAT5 that skipped exon 3, exon 9, or exon 10; another variant skipped a large portion of exon 7, while a fifth variant skipped part of exon 8 and all of exon 9 (Fig. 3). Each of the EAAT5 variants was shown to be expressed at the protein level by Western blotting using splice site–specific antibodies (Lee et al. 2012a). EAAT5 is particularly interesting since this protein exhibits a relatively large chloride conductance, associated with modest transport activity, suggesting a function more closely related to ligand-gated chloride channels than classic transporters. This has led to the finding that wild-type EAAT5 functions as an "inhibitory" presynaptic glutamate receptor in retinal bipolar cells (Wersinger et al. 2006).



Currently, it is unknown whether the EAAT5 splice variants could function as transporters or receptors. These variants appear to be missing key structural features that may be important for chloride conductance or glutamate binding/ transport. Thus, the functional characterization of such variants may require analysis beyond the determination of whether they conduct chloride or translocate glutamate.

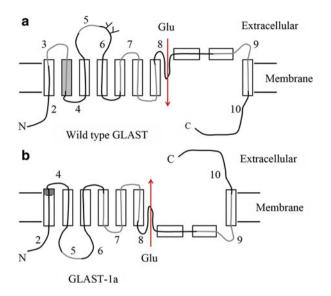
# 2.6 Splicing of N- and C-Termini

At least, three 5'-variants of GLT-1 have been identified in brain which contain varying lengths of upstream open-reading frames encoding GLT-1 proteins with distinct N-terminal sequences: MASTEGANN, MPK, or MKRPKEHSIQRS (Rozyczka and Engele 2005; Münch et al. 2002; Peacey et al. 2009). A different 5'-variant of GLT-1 has also been identified in liver which contain the N-terminal sequence MVS (Utsunomiya-Tate et al. 1997); differences in the 5'-ends of the GLT-1 transcripts in brain and liver suggest the presence of tissue-specific promoters. In addition, at the 3' end of the GLT-1 gene, there is extensive differential exon usage that can result in the generation of at least three distinct C-termini sequences (see Fig. 2b). Thus, at least 12 variant forms of GLT-1 protein potentially exist, depending on the splicing at the 5' and 3'-ends of the GLT-1-coding sequence. Uptake studies in cultured cells and Xenopus oocytes have shown that functional and pharmacological properties of GLT-1 variants are not changed by alterations in their N- and C-termini sequences (Utsunomiya-Tate et al. 1997; Peacey et al. 2009). Therefore, N- and C-terminal splicing may be more important for differential expression or targeting of the transporters to particular cell types than in regulating the properties of glutamate uptake.

# 3 Are the EAAT Variants Generated by Alternative Splicing Functional?

While N- and C-terminal splicing can produce functional EAAT variants such as GLT-1b and GLT-1c, very little information is available on the functional properties of exon-skipping variants such as GLAST-1a, GLAST-1b, and GLAST-1c. The exon 9-skipping GLAST-1b has received little attention because when overexpressed in HEK293 cells, it was localized to the endoplasmic reticulum (ER) and thus is unable to act as a plasmalemmal glutamate transporter. When co-expressed with wild-type GLAST, it exerted a dominant negative effect over GLAST function and reduced GLAST insertion into the plasma membrane, suggesting that GLAST-1b could potentially act as an interfering subunit of the GLAST homo-oligomeric structure (Vallejo-Illarramendi et al. 2005). However, when a specific antibody was later developed to GLAST-1b (Macnab and Pow 2007b), it was shown that rather than being restricted to the ER, GLAST-1b was readily demonstrable in the plasma membranes of a very small subset of neurons. Not only was GLAST-1b targeted to the plasma membrane, which would be essential for its role as a transporter but it was also shown that the expression levels rise dramatically in neurons after a hypoxic insult, indicating that expression is physiologically regulated (Sullivan et al. 2007a). Similar issues have arisen for GLAST-1a, which is an exon 3-skipping variant of GLAST. Interestingly, some algorithms have predicted that the loss of exon 3 would produce a reversal of the membrane topology of GLAST-1a (Huggett et al. 2000), where the orientation of the glutamate-binding/-translocating region and the direction of glutamate transport may also be reversed (Fig. 4). This does not however imply a loss of function but rather may indicate a different function for GLAST-1a, such as acting as a glutamate "efflux" protein. Patch clamping experiments in brain slices have clearly demonstrated that glutamate transporters can operate in reverse, transporting glutamate from inside to outside of the cell under conditions such as anoxia (Billups and Attwell 1996; Rossi et al. 2000). Furthermore, alternative splicing of the bile acid transporter (ASBT) gene is known to generate an exon 2-skipping variant that functions as a bile acid efflux protein and localized to basolateral domains of cholangiocytes. This is in contrast to wild-type ASBT which functions to take up bile acids across the apical membranes of cholangiocytes (Lazaridis et al. 2000). Thus, alternative splicing can alter the cellular targeting and/or functional properties of transporters.

In the case of GLAST-1c, which skips exons 5 and 6, this variant has significant key features in common with the prototypical bacterial aspartate transporter GLTph (Yernool et al. 2004), which appears to be an ancestral form of glutamate transporter; these include the absence of glycosylation sites, and much of the same extracellular loop that is missing in GLAST-1c is also absent in GLTph. Such similarities between GLAST-1c and a functional prototypic transporter may support the view that GLAST-1c is plausibly a functional glutamate transporter and possibly represents a primitive form of GLAST. Accordingly, the possibility that GLAST-1a, GLAST-1b, and GLAST-1c are involved in either directly transporting glutamate or in regulating the trafficking of wild-type GLAST must remain a possibility at this stage.



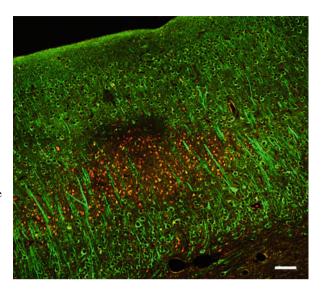
**Fig. 4** Predicted transmembrane (TM) topologies of wild-type GLAST and GLAST-1a; the extracellular surface and the plasma membrane are indicated. Exon boundaries are overlaid onto the predicted topologies to demonstrate the predicted changes in topology that result from the skipping of exon 3. Deletion of exon 3 in GLAST-1a converts the first two TM domains into one and is predicted to reverse the membrane topology, resulting in an extracellular C-terminal region that may be associated with the reversal of glutamate transport (Adapted from Huggett et al. 2000)

# 4 Disease Association of EAAT Splice Variants

Much evidence exists for the differential regulation of EAAT2 and EAAT2b in amyotrophic lateral sclerosis (ALS) Meyer et al., 1999. Dramatic reduction in EAAT2 protein has been observed in motor cortex of ALS patients without alterations in EAAT2 (GLT-1) mRNA levels, suggesting posttranscriptional regulation may be involved (Bristol and Rothstein1996; Rothstein et al. 1995; Fray et al. 1998). This is in contrast to expression of EAAT2b (GLT-1b) protein, which is increased in motor cortex of ALS patients (Maragakis et al. 2004). A number of animal models have been used to study the potential role(s) of EAAT expression in ALS. Studies in mice where the animals are fed with cycads reveal behavior and histological features resembling ALS-PDS. In these animals, there is a significant downregulation of GLT-1a protein in spinal cord and primary somatosensory cortex; GLT-1b protein levels were significantly decreased in the spinal cord, in the motor, somatosensory, and piriform cortices, and in the striatum. Notably, GLT-1b expression was decreased in areas where GLT-1a expression was unaltered, suggesting differential regulation of the two transporter subtypes (Wilson et al. 2003). Similarly, Munch et al. (2002) have also shown that the N-terminal splicing of GLT-1a was altered in the spinal cord of SOD transgenic mice, suggesting a link between altered splicing of the 5'-end sequences of GLT-1

gene and ALS. Whether changes in expression or splicing of GLT-1 are aspects of early causation, or a consequence of ALS, remains open to interpretation. However, Guo et al. (2003) produced a double transgenic SOD mutant mice overexpressing GLT-1a which showed significant delays in grip-strength decline and a delay in the loss of motor neurons (Guo et al. 2003). However, these transgenic animals did not show decline in the onset of paralysis nor did they live longer. These results suggest that loss of GLT-1/EAAT2 may contribute to, but do not initially cause, motor neuron degeneration in ALS (Guo et al. 2003).

Aberrant expression of EAAT splice variants has been observed in Alzheimer's disease (AD). Scott et al. (2011) showed increased mRNA expression of an exon 9-skipping form of EAAT2 and protein for this variant has been observed in neurons of AD brain (Pow and Cook 2009). Similarly, protein for the exon 3-skipping form of EAAT1 has been observed in neurons of the AD cortex (Pow and Cook 2009). It is plausible that changes to the splicing of GLT-1 (EAAT2) and GLAST (EAAT1) may explain why there has been a paradoxical observation of an apparently decreased GLAST and GLT-1 expression but increased glutamate reuptake activity in cortex of ABPP23 mice (Schallier et al. 2011), suggesting possibly that modified forms of the transporters were present and functional but not detected with the tools used. Aberrant neuronal expression of EAAT splice variants has also been observed in other pathophysiological states. Thus, Sullivan et al. (2007a) have shown expression of an exon 9-skipping form of GLAST (GLAST-1b) in populations of neurons after a hypoxic insult. GLT-1b, which has an alternatively spliced C-terminal region, was also shown to be aberrantly expressed in neurons after hypoxic insult (Pow et al. 2004). In both scenarios, it has been proposed that an elevation of extracellular glutamate concentration may be driving the expression of the EAAT subtypes. Similar observations have been seen in disease states such as glaucoma and age-related macular degeneration (AMD) as well as in stroke where glutamate excitotoxicity has been implicated. For example, induced expression of a C-terminal splice variant of GLT-1 (GLT-1c) has been demonstrated in retinal ganglion cells in response to elevated intraocular pressure in rats or in human eyes afflicted by glaucoma (Sullivan et al. 2006), and this has been interpreted as being adaptive, to protect the ganglion cells that are vulnerable to glutamate excitotoxicity. Conversely, a complete loss of GLT-1c from photoreceptors in peri-macular regions was observed in AMD patients (Pow et al. 2005) and may explain why these cells lose their content of glutamate that acts both as a neurotransmitter and as an essential metabolic fuel for these cells, which subsequently die. In stroke, induced expression of the exon 3-skipping form of GLAST (GLAST-1a) has been observed in cerebral cortex of ischemic rats induced by middle cerebral artery occlusion (MCAo) (Fig. 5; Pow et al. Unpublished data). Thus, anomalies in local excitation (as a result of ischemic stroke) can induce expression of glial glutamate transporter splice variants such as GLAST-1a in affected neurons, possibly as a protective mechanism. This is in contrast to expression of GLT-1a mRNA and protein, which was reported to be decreased in cerebral cortex of ischemic rats induced by MCAo (Rao et al. 2001a). Interestingly, decreasing GLT-1a expression by antisense knockdown was shown to **Fig. 5** Image of rat cortex after a stroke induced by transient middle cerebral artery occlusion, immunolabeled for MAP2 (*green*) which labels cell bodies and dendrites of healthy neurons. The large central area where the neurons have lost MAP2 labeling represents the incipient lesion area; GLAST-1a labeling (*red*) is upregulated in this area. Scale bar, 50 µm



exacerbate neuronal damage induced by MCAo (Rao et al. 2001b) while targeted overexpression of GLT-1a reduced ischemic brain injury caused by MCAo (Harvey et al. 2011), suggesting that increasing the capacity to remove extracellular glutamate may provide beneficial outcomes against ischemia-induced glutamate excitotoxicity.

In Schizophrenia, an ambiguous literature exists concerning the expression level of EAATs with increases and decreases being reported. Thus, Smith et al. (2001) examined postmortem tissues from Schizophrenia subjects by in situ hybridization and showed elevated expression of EAAT2 mRNA in thalamus. Similar studies by Ohnuma et al. (1998, 2000) have reported reduced expression of EAAT2 mRNA in prefrontal cortex and parahippocampal regions of Schizophrenia subjects. Conversely, Bauer et al. (2008) did not detect changes in transcript or protein expression of EAAT2 in postmortem prefrontal cortex tissues of Schizophrenia patients using in situ hybridization and Western blotting. One possible limitation of these studies is the putative effects of psychotropic medications on EAAT expression. Several studies have addressed this issue by examining expression of EAAT2 (GLT-1) in animals treated with such drugs. Melone et al. (2001) have shown that short-term (3 weeks) administration of oral clozapine (25–35 mg/kg per day) to rats reduced expression of GLT-1a in cerebral cortex, resulting in decreased glutamate transport and increased extracellular glutamate concentration. Suppression of GLT-1a expression was also observed in rat brain after long-term (6 months) chronic antipsychotic treatment with clozapine (45 mg/kg per day) (Schmitt et al. 2003). The current literature, therefore, suggests that GLT-1a expression is regulated by exposure to antipsychotic drugs. There is no data at present as to whether changes in GLT-1a expression are due to changes in splicing or to changes in overall transcription or translation of the mRNA.

#### 5 Conclusion

Alternative splicing of EAATs is a common mechanism that allows for diversification of a single gene into different protein products. For EAAT2, many of the splice variants produced by N- and C-terminal splicing have been shown to be functional, however, with changes in tissue distribution. For other EAAT members such as EAAT1 and EAAT5, "exon skipping" is the most common mode of splicing whereby certain exons have been removed, therefore resulting in shorter protein products. This highly versatile mechanism can regulate the amount of mRNA and changes in protein sequence that is likely to affect many aspects of protein function such as substrate-binding properties, posttranslational modifications, cellular localization, and trafficking. Interestingly, some of the "exon skipping" EAAT variants cloned to date lack significant portions of the C-terminal regions, suggesting that critical interactions such as binding to the cytoskeleton of the cell would be differentially affected in such variants. Conversely, others lack portions of the N-terminal region and thus may exhibit deficiencies in parameters such as chloride conductances. Where key structural elements of a transporter are missing, it is often dismissed as "junk protein," but biological systems are extremely conservative and unlikely to waste energy making junk in physiological situations. Instead we suggest, by homology to more recent ideas on "junk DNA," that it is plausible that many of these novel proteins may instead have primary functions that are distinct from their initially described roles as glutamate transporters or receptors. In some cases, the splice variants have been shown to possess a dominant negative effect on the co-transporter function, suggesting an important role in the functional regulation of the wild-type transporter. Validating the function of such EAAT variants will require further experimental analysis. From a clinical perspective, however, these EAAT variants are of particular interest, given that their expression and/or regulatory pathways are associated with neurological disorders such as Alzheimer's disease, glaucoma, and macular degeneration as well as hypoxia. Determining whether EAAT changes in any of these disease states are causal or consequential and whether modulation of the activity for a specific EAAT variant might evolve as a therapeutic option are just two examples for the many questions waiting to be answered. Further studies of alternative splicing regulation of EAATs will no doubt reveal new opportunities.

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# AIDS

# M. Kaul

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#### Abstract

Infection with human immunodeficiency virus (HIV)-1 not only targets and compromises cells of the immune system and eventually causes AIDS but also triggers a variety of neurological problems that can culminate in frank dementia. Several lines of evidence suggest that HIV-1 strikes at the brain in at least two ways: (i) by causing toxicity leading to neuronal injury and death and (ii) by impairing neurogenesis without killing neural precursors. Both of the pathogenic mechanisms share the involvement of neuroinflammatory mediators, and both processes seem to be influenced by host and viral factors, including the interferon response, age and viral subtypes, as well as treatment and comorbidity factors, such as drug abuse. Development of both HIV-associated neurocognitive disorders (HAND) and AIDS goes hand-in-hand, and despite

M. Kaul

Infectious and Inflammatory Disease Center, Sanford-Burnham Medical Research Institute, La Jolla, CA, USA

e-mail: mkaul@sanfordburnham.org; mkaul@sbmri.org

viral control and life-prolonging effects of combination antiretroviral treatments (cART), neuroinflammation and concomitant activation of macrophages and microglia accompany the continued development of HAND in HIV patients. Besides intact HIV-1, six of its proteins have been linked to neuronal injury. Experimental evidence supports two different, but not exclusive, hypotheses of how HIV-1 causes neuronal damage in the brain. The "direct injury" hypothesis proposes that viral proteins directly compromise neurons, whereas the "indirect" or "bystander effect" hypothesis states virus infection and viral components trigger in macrophages and microglia and astrocytes the production of neurotoxins that injure neurons. HIV-1 and its six neurotoxic protein components can trigger a plethora of cellular responses and signaling mechanisms. However, it appears that the HIV-1 and its viral components cause neurotoxicity via converging mechanisms even if the virus and each component may start out their pathogenic mechanism by interacting with different host cell factors. Converging mechanisms in HIV neurotoxicity include neuroinflammation, MAPK signaling, altered gene expression, oxidative stress, interference with the cell cycle machinery and autophagy, excitotoxicity, synaptic injury, and, ultimately, neuronal apoptosis.

#### Keywords

HIV-1 • Infection • Inflammation • Brain • Cognition • Dementia • Neurotoxicity
• Excitotoxicity • Neurogenesis • Neurons • Astrocytes • Macrophages • Microglia • Host-pathogen Interaction • Immune Response • Co-morbidity • Drug Abuse

List of Abbreviati	ions
AIDS	Acquired immunodeficiency syndrome
ANI	Asymptomatic neurocognitive impairment
ANT	Adenine nucleotide translocator
ApoE3	Apolipoprotein E3
BBB	Blood-brain barrier
cART	Combination antiretroviral therapy
CEBP	CCAAT-enhancer-binding protein
CNS	Central nervous system
COUP-TF	Chicken ovalbumin upstream promoter transcription factor
CSF	Cerebrospinal fluid
Dox	Doxycycline
EAAs	Excitatory amino acids
Env	Envelope protein
ER	Endoplasmic reticulum
ERK-1/ERK-2	Extracellularly regulated kinase-1 and kinase-2
GABA	γ-aminobutyric acid

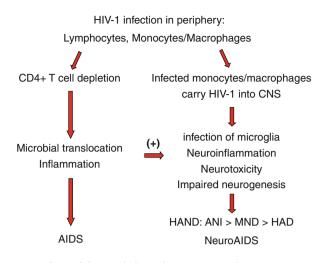
GSK	Glucagon synthase kinase
HAART	Highly active antiretroviral therapy
HAD	HIV-associated dementia
HAND	HIV-associated neurocognitive disorders
HIV-1	Human immunodeficiency virus-1
HIVE	HIV encephalitis
HNE	Hydroxynonenal
IFN	Interferon
IL	Interleukin
iNOS	Inducible nitric oxide synthase
IRIS	Immune reconstitution inflammatory syndrome
LPS	Lipopolysaccharide
LRP	Low-density lipoprotein receptor-related protein
LTP	Long-term potentiation
MAPK	Mitogen-activated protein kinase
MCMD	Minor cognitive motor disorder
MCP	Monocyte chemoattractant protein
mGluRs	Metabotropic glutamate receptors
miR	MicroRNAs
MLK	Mixed lineage kinase
MMPs	Metalloproteinases
MND	Mild neurocognitive disorder
Nef	Negative factor
NMDAR	<i>N</i> -methyl- <i>D</i> -aspartate-type glutamate receptor
Pol	Polymerase
PTPC	Permeability transition pore complex
Rb	Retinoblastoma protein
Rev	Regulatory for expression of viral proteins
RyR	Ryanodine receptors
SCID	Severe combined immunodeficiency
SIV	Simian immunodeficiency virus
Tat	Transactivator of transcription
tg	Transgenic
TLR	Toll-like receptors
TNF	Tumor necrosis factor
TNFR	TNFa receptor
TRAIL	TNF-related apoptosis-inducing ligand
TUNEL	Terminal-deoxynucleotidyl-transferase-mediated dUTP nick-end
	labeling
UPR	Unfolded protein response
Vif	Viral infectivity factor
Vpr	Viral protein r
Vpu	Viral protein u

#### 1 Introduction

Human immunodeficiency virus-1 (HIV-1) is a retrovirus belonging to the subgroup of lentiviruses (Barre-Sinoussi et al. 1983; Alizon et al. 1984; Hahn et al. 1984; Wain-Hobson et al. 1985). HIV was discovered in 1983 using lymphoid tissue from one of a quickly increasing number of patients presenting with lymphadenopathy followed by an acquired immunodeficiency syndrome (AIDS) (Barre-Sinoussi et al. 1983; Hahn et al. 1984; Barre-Sinoussi 2003). In 2011, an estimated 34.2 million people were living with HIV infection or AIDS, and despite improvements in therapy and access to treatments, the epidemic caused 2.5 million deaths (UNAIDS; www.unaids.org).

Although HIV-1 primarily targets and compromises cells of the immune system and eventually causes AIDS (Stevenson 2003), it was observed early on in the epidemic that the virus can also cause a variety of neurological problems that can culminate in frank dementia (Loewenstein and Sharfstein 1983; Jordan et al. 1985; Petito et al. 1985). While AIDS-related opportunistic infections obviously affected the central nervous system (CNS), in particular in the absence of treatment, HIV-1 infection itself seemed also able to initiate a number of neurological syndromes (Jordan et al. 1985; Petito et al. 1985). Today it is widely accepted that HIV-1 can directly trigger neurological conditions, including peripheral neuropathies, vacuolar myelopathy, and a clinical syndrome of cognitive and motor dysfunction that has early on been designated AIDS dementia and later on HIV-associated dementia (HAD) (Navia et al. 1986a, b; Petito et al. 1986). A milder and, in the presence of antiviral treatment, more common form of HAD is termed minor cognitive motor disorder (MCMD) (Ellis et al. 1997; Kaul et al. 2001; Gendelman et al. 2005; Power et al. 2009). In recognition of a changing clinical presentation, neurocognitive complications of HIV-1 infection have more recently been categorized as HIVassociated neurocognitive disorders (HAND) (Antinori et al. 2007). HAND defines three categories of disorders with increasing severity according to standardized measures of dysfunction: (i) asymptomatic neurocognitive impairment (ANI), (ii) mild neurocognitive disorder (MND), and (iii) HAD.

The mechanisms contributing to the development of the various forms of HAND remain incompletely understood, although it is known that anemia early in HIV-1 infection appears to presage later neuropsychological impairment (McArthur et al. 1993). In order to learn how HIV infection can lead to dementia or the milder forms of HAND, scientists have employed a variety of strategies. Besides neurological and neuropsychological assessments, which are essential to identify the clinical domains of HAND, the current knowledge is based on multiple methodological approaches, including analyses of cerebrospinal fluid (CSF), neuroimaging, histopathological studies on *postmortem* brain specimens, animal models ranging from monkeys to rodents, and in vitro models using neuronal, glial, and immune cells from humans, monkeys, and rodents (Maung et al. 2012). New and sometimes surprising insights into the ways HIV-1 compromises the CNS came from the



**Fig. 1** Schematic outline of key pathological processes and events associated with HIV-1 infection and the development of HAND, neuroAIDS, and AIDS. HIV-1 infection occurs in the periphery, but soon afterwards the virus arrives in the brain, presumably carried by infected infiltrating macrophages and possibly lymphocytes. In the brain, the virus can infect microglia. Infected macrophages and microglia release viral proteins and host factors that can stimulate uninfected cells contributing to the development of neuroinflammation, neurotoxicity, and impaired neurogenesis. The latter two processes are presumed to underlie the development of HAND. In the periphery, HIV infection most often leads to the loss and eventual depletion of  $CD4^+ T$  cells, including those in the intestinal tract. The weakened immune surveillance seems to enable microbial translocation, which further contributes to inflammation. Depletion of  $CD4^+ T$  cells will eventually result in the development of AIDS. Peripheral inflammation also seems to promote neuroinflammation and the development of HAND and neuroAIDS

discovery of HIV-1 coreceptors, the chemokine receptors CCR5 and CXCR4, in the brain as well as from progress in understanding neuroinflammation and neural stem cell biology (Lavi et al. 1998; Miller and Meucci 1999; Kaul et al. 2001; Kramer-Hammerle et al. 2005; Minghetti 2005; Kaul et al. 2005; Gonzalez-Scarano and Martin-Garcia 2005; Jones and Power 2006). Currently, there is strong evidence suggesting that HIV-1 may strike at the brain in at least two ways: (i) by causing toxicity leading to neuronal injury and death and (ii) by impairing neurogenesis without killing neural precursors (Fig. 1) (Kaul 2008). Both of the two mechanisms appear to share the involvement of inflammatory processes, and both seem to be influenced by host and viral factors, such as drug abuse (Kaul 2008; Kaul 2009; Gannon et al. 2011). While we cannot yet reliably determine how much impairment of neurogenesis contributes to HAND, based on currently available information, this chapter will discuss what may be the most destructive mechanism for the CNS during HIV-1 infection: neurotoxicity.

# 2 HIV-1 Infection, AIDS, and HIV-Associated Neurocognitive Disorders

HAD as the most severe manifestation of HAND occurred at the beginning of the AIDS epidemic primarily in patients with advanced HIV disease and CD4<sup>+</sup> T cell counts below 200  $\mu$ l<sup>-1</sup> (McArthur et al. 1993). The introduction in the mid-1990 of combination antiretroviral therapy (cART), which was also called "highly active antiretroviral therapy" (HAART), was a major advance in the treatment of HIV infection that in most cases significantly delayed the progression to AIDS. Thus, cART increased the life span of people infected with HIV-1 and resulted in a temporary decrease in the incidence of HAD from 20 % to 30 % to as low as 10.5 % (McArthur et al. 2003). Control of peripheral viral replication and the treatment of opportunistic infections continue to extend survival times, and HIV infection has become a chronic rather than acute disease. Unfortunately, however, cART fails so far to prevent the eventual development of MND or HAD or to reverse HAND (Cunningham et al. 2000; Cysique et al. 2006; Giancola et al. 2006; Nath and Sacktor 2006; Ellis et al. 1997; Antinori et al. 2007; Liner et al. 2008; Boisse et al. 2008; Brew et al. 2008; Heaton et al. 2010, 2011; Gannon et al. 2011). In the era of cART, MND may be more prevalent than overt dementia, but as an AIDS-defining illness, the incidence of dementia has increased in recent years and HAD still constitutes a significant independent risk factor for death due to AIDS (Ellis et al. 1997; Liner et al. 2008). In a group of 669 HIV patients who died between 1996 and 2001, more than 90 % had been diagnosed with HAD as an AIDS-defining condition within the last 12 months of life (Welch and Morse 2002). In addition, the proportion of new cases of HAD displaying a CD4<sup>+</sup> T cell count greater than 200  $\mu$ l<sup>-1</sup> has been growing (McArthur et al. 2003). Of note, recent therapeutic guidelines for AIDS suggest to start cART only once the number of CD4<sup>+</sup> T cells begins to decline. Since this might occur up to years after initial peripheral infection, cART most likely has no chance to forestall the entry of HIV-1 into the CNS (Kramer-Hammerle et al. 2005). Furthermore, distinct patterns of viral drug resistance in plasma and CSF compartments have been ascribed to limited penetration of certain antiretroviral drugs, such as HIV protease inhibitors and several of the nucleoside analogues, into the CNS (Cunningham et al. 2000; Kaul et al. 2005; Kramer-Hammerle et al. 2005). On the other hand, several studies have found clear evidence that cART can benefit the brain and neurocognitive performance, in particular if the therapeutic compounds reach the CNS in concentrations sufficient to suppress virus (Smurzynski et al. 2011; Letendre et al. 2004; Patel et al. 2009; Winston et al. 2010; Gannon et al. 2011). At the same time, some studies suggest that long-term exposure to cART poses a potential toxicological problem, which may contribute to neurocognitive deterioration on its own (Liner et al. 2008, 2010; Letendre et al. 2008; Cardenas et al. 2009; Marra et al. 2009; Schweinsburg et al. 2005).

Early studies during the AIDS epidemic indicated that HIV penetrates into the CNS soon after infection in the periphery and then resides primarily in perivascular macrophages and microglia (Ho et al. 1985; Gartner 2000; Gonzalez-Scarano and Martin-Garcia 2005). Therefore it is not very surprising that despite cART the

prevalence of dementia may be rising as people live longer and age with HIV-1 infection (Lipton 1997; Jones and Power 2006; Cunningham et al. 2000; McArthur et al. 2003; Kaul et al. 2005, 2001; Kramer-Hammerle et al. 2005; Brew et al. 2009; Lindl et al. 2010; Gannon et al. 2011).

The blood–brain barrier (BBB) seems to play an important role in the acute and chronic HIV infection of the CNS as it potentially controls the infiltration of virusinfected and uninfected peripheral immune cells, such as monocytes and macrophages (Gartner 2000; Nottet et al. 1996; Gras and Kaul 2010). Astrocytes and microglia produce chemokines – cell migration-/chemotaxis-inducing cytokines – such as monocyte chemoattractant protein (MCP)-1, which appear to attract peripheral blood mononuclear cells across the BBB into the brain parenchyma (Asensio and Campbell 1999). In fact, an increased risk of HAD has recently been connected to a mutant MCP-1 allele that causes increased infiltration of monocytes and macrophages into tissues which may enhance the transfer of HIV-1 into the brain and promote neuroinflammation (Gonzalez et al. 2002; Gras and Kaul 2010). Alternatively, it has been suggested that the proinflammatory cytokine TNFα promotes a paracellular route for free virus across the BBB (Fiala et al. 1997).

Importantly, infectious events in the periphery and the associated immune response and inflammatory processes influence all cell types in the CNS (Turrin and Rivest 2004; Chakravarty and Herkenham 2005). Potentially important for the understanding of all aspects of HIV disease are recent findings strongly suggesting that the pathogenic or nonpathogenic course of HIV/simian immunodeficiency virus (SIV) infection may be determined by the reaction of plasmacytoid dendritic cells to the virus, in particular the time and amount of interferon (IFN)- $\alpha$  production (Mandl et al. 2008). Signaling via Toll-like receptors (TLR)-7 and -9 and the extent of downstream activation of interferon regulatory factor (IRF)-7 appears to control how much IFN $\alpha$  is being produced. While IFNs are important for an antiviral immune response, the lasting production of IFN $\alpha$  in HIV/SIV infection is suspected to result in erroneous and exhaustive immune activation which in turn may lead to immune suppression and progression to SIV/AIDS (Mandl et al. 2008). Besides immune suppression, the extensive production of IFN $\alpha$  can be of direct detrimental consequence to the brain, as IFN production in the HIV-infected CNS directly correlates with neurocognitive impairment (Sas et al. 2007). In contrast, IFNβ induces a dominant-negative form of the transcription factor CCAATenhancer-binding protein (CEBP)- $\beta$  and suppresses SIV in macaque macrophages (Dudaronek et al. 2007). Moreover, IFNβ triggers in macrophages and microglia the expression of the antiviral and neuroprotective  $\beta$ -chemokines CCL3/MIP-1 $\alpha$ , CCL4/MIP-1β, and CCL5/RANTES (Kitai et al. 2000; Garzino-Demo et al. 1998; Kaul and Lipton 1999; Kaul et al. 2007).

HIV-1/SIV infection also affects the intestinal tract and can cause leakage of bacteria or bacterial products into the bloodstream (Brenchley et al. 2006). Such microbial translocation is reflected by elevated plasma levels of lipopolysaccharide (LPS), which is associated with increased monocyte activation and dementia in HIV-infected patients (Ancuta et al. 2008). Thus, immune activation and inflammation are prime suspects regarding a contribution to neurotoxicity, impairment of

neurogenesis, and the development of HAND, but the biological effects of IFN $\beta$  suggest that innate immunity can also protect.

# 3 Neuropathology of HIV-1 Infection Before and After Introduction of cART

Development of HAND and in particular HAD in patients during their lifetime has been correlated to a *postmortem* neuropathological diagnosis that has often been referred to as HIV encephalitis (HIVE) (Petito et al. 1986; Masliah et al. 1992). The salient pathological features include microglial nodules; activated resident microglia; HIV-infected multinucleated giant cells; infiltration predominantly by monocytoid cells, including blood-derived macrophages; widespread reactive astrocytosis; myelin pallor; and decreased synaptic and dendritic density, combined with selective neuronal loss (Petito et al. 1986; Masliah et al. 1997). The designation and hallmarks of HIVE indicate that signs of neuroinflammation have commonly been found in HIV patients since the beginning of the AIDS epidemic. Surprisingly, however, antemortem measures of cognitive dysfunction did not correlate well with numbers of HIVinfected cells, multinucleated giant cells, or viral antigens in CNS tissue (Glass et al. 1995; Achim et al. 1994; Wiley et al. 1994; Masliah et al. 1997). Instead, the pathological characteristics showing the best correlation with clinical signs of HAD were increased numbers of microglia (Glass et al. 1995), decreased synaptic and dendritic density, selective neuronal loss (Masliah et al. 1997; Achim et al. 1994; Wiley et al. 1994), elevated tumor necrosis factor (TNF)- $\alpha$  mRNA in microglia and astrocytes (Wesselingh et al. 1997), and evidence of excitatory neurotoxins in CSF and serum (Heyes et al. 1991). Thus, several lines of evidence pointed toward an important role of inflammatory factors, macrophages/microglia, and neurotoxins in the development of HAD. In addition, two recent studies also suggested a crucial role of mononuclear phagocytes because the findings showed that the amount of proviral HIV DNA in circulating monocytes and macrophages correlated well with HAD, whereas viral load was a less reliable indicator (Shiramizu et al. 2005, 2006).

Neuronal damage and loss has been observed *postmortem* in distinct brain regions of HIV and AIDS patients, including frontal cortex (Ketzler et al. 1990; Everall et al. 1991), substantia nigra (Reyes et al. 1991), cerebellum (Graus et al. 1990), and putamen (Everall et al. 1993). Focal neuronal necrosis appeared to occur in HAD brains early in the AIDS epidemic (Ho et al. 1989), whereas neuronal apoptosis has more recently been linked to HAD (Petito and Roberts 1995; Adle-Biassette et al. 1999). The localization of apoptotic neurons was correlated with signs of structural atrophy and microglial activation, in particular within subcortical deep gray structures (Adle-Biassette et al. 1999). Surprisingly, however, neuronal death was not clearly associated with viral burden or a history of dementia. On the other hand, a separate study found that cellular DNA by terminal-deoxynucleotidyl-transferase-mediated fragmentation detected dUTP nick-end labeling (TUNEL) in basal ganglia correlated well with HAD (Rostasy et al. 2000).

To be sure, the neuropathology of HIV infection and AIDS has shifted following the introduction of cART (Anthony and Bell 2008; Boisse et al. 2008; Brew et al. 2008). Although the number of opportunistic infections declined, two *postmortem* studies found evidence of increased infiltration and activation of macrophages and microglia in hippocampus and basal ganglia of cART-experienced patients in comparison to cases from the time before cART and a higher prevalence of HIVassociated encephalitis at the time of autopsy (Anthony et al. 2005; Bell 2004; Langford et al. 2003). Even more encephalitis and severe leukoencephalopathy was observed in *postmortem* specimens from AIDS patients who had failed cART (Anthony et al. 2005). In line with these findings, more recent neuropathological descriptions also reported various forms with severe HIVE and white matter injury, extensive perivascular lymphocytic infiltration, "burnt-out" forms of HIVE, and aging-associated  $\beta$ -amyloid accumulation with Alzheimer's-like neuropathology (Everall et al. 2005; Green et al. 2005; Achim et al. 2009).

Signs of neuroinflammation were commonly found in HIV patients since the beginning of the AIDS epidemic (HIVE) (Masliah et al. 1992), and neuroinflammation usually developed with the progression of HIV patients from the latent, asymptomatic stage of the disease to AIDS and HAD. Therefore, inflammation itself was generally seen as a pathological mechanism, and improved therapy was expected to ameliorate neuroinflammation. Surprisingly, autopsy cases of HIV-related death collected since the advent of cART rather suggested the opposite (Bell 2004; Everall et al. 2005; Anthony et al. 2005). However, the predominant localization of neuroinflammation had changed. Specimen from pre-cART patients showed strong involvement of basal ganglia, whereas post-cART cases displayed extensive signs of inflammation in hippocampus and adjacent regions of entorhinal and temporal cortex (Anthony and Bell 2008). Interestingly, cART seemed to have limited or prevented lymphocyte infiltration, with the exception of occasional, distinct instances, termed "immune reconstitution inflammatory syndrome" (IRIS), where massive lymphocytosis was associated with pronounced demyelination and white matter damage (Anthony and Bell 2008; Boisse et al. 2008). Thus, despite viral control and life-prolonging effects of cART, neuroinflammation and concomitant activation of macrophages and microglia accompany the continued development of HAND in HIV patients.

# 4 Chemokine Receptors and HIV-1 Infection of the Brain

Chemokine receptors are members of the large family of G-protein-coupled seven transmembrane-spanning domain receptors. Their ligands are primarily small proteins, the chemokines, which trigger leukocyte trafficking and participate intimately in the inflammatory and anti-infection responses of the immune system (Locati and Murphy 1999; Dragic et al. 1996; Oberlin et al. 1996; Tran and Miller 2003). Beyond the immune defense, chemokines and their receptors contribute to organogenesis, including hematopoiesis, angiogenesis, and development of heart and brain (Tachibana et al. 1998; Zou et al. 1998; Ma et al. 1998; Locati and Murphy 1999), and are essential for maintenance, maturation, and migration of hematopoietic and neural stem cells (Lapidot and Petit 2002; Tran and Miller 2003). However, some pathogens are able to hijack certain chemokine receptors for their purposes, such as the mediation of HIV-1 infection (Dragic et al. 1996; Oberlin et al. 1996; Locati and Murphy 1999; Sodhi et al. 2004).

HIV-1 infects macrophages and lymphocytes in the periphery and microglia in the brain after binding of the viral envelope protein gp120 to CD4 in conjunction with one of several possible chemokine receptors. CD4 and HIV-1/gp120 coreceptors are likely the first sites of host-virus interaction. Different HIV-1 strains may use CCR5 (CD195) - and CCR3 on microglia - or CXCR4 (CD184) or a combination of these chemokine receptors to enter target cells, depending on specific properties of their gp120 (Dragic et al. 1996; Oberlin et al. 1996; He et al. 1997a). Generally, "T-tropic" viruses predominantly infect T lymphocytes via the  $\alpha$ -chemokine receptor CXCR4 and/or the  $\beta$ -chemokine receptor CCR5. In contrast, "M-tropic" HIV-1 primarily enter macrophages and microglia via CCR5 and CCR3, although CXCR4 may also be involved (He et al. 1997a; Michael and Moore 1999; Ohagen et al. 1999; Chen et al. 2002). Of note, usage of CCR5 is neither necessary nor sufficient for macrophage tropism, and CXCR4-preferring viruses can infect macrophages (Gorry et al. 2005; Li et al. 1999; Yi et al. 1998, 2003). In any case, chemokine receptors play crucial roles on several levels in HIV-1 disease (Verani and Lusso 2002). Usually, CCR5preferring HIV-1 strains (R5-tropic) are transmitted between humans, and individuals lacking CCR5 are highly resistant to primary HIV infection (Liu et al. 1996). CXCR4-using viruses (X4-tropic) occur in about 50 % of infected individuals later in the course of HIV-1 disease and generally indicate progression to AIDS (Michael and Moore 1999). These X4-tropic viruses can easily be transmitted via blood transfusion or shared needles in intravenous drug users. However, in many HIV/AIDS patients, R5-tropic viruses evolve over time into more cytopathic variants possessing higher CCR5 affinity combined with diminished CD4 dependence, and a switch in coreceptor usage does not occur (Gorry et al. 2005).

In the brain, neurons and astrocytes express among other chemokine receptors also CCR5 and CXCR4 (Asensio and Campbell 1999; Miller and Meucci 1999), although these cell types, in contrast to microglia, are not thought to harbor productive HIV-1 infection under in vivo conditions. However, HIV infection of astrocytes seems to occur and require the help of a mannose receptor (Liu et al. 2004). Contrasting earlier reports indicating nonproductive infection, recent studies suggest that under certain conditions of immune activation, astrocytes may be able to produce virus (Li et al. 2011). In any case, several in vitro studies strongly suggest that CXCR4 is prominently involved in HIV-associated neuronal damage, whereas CCR5 may play a dual role by mediating either a toxic or protective effect depending on the kind and concentration of its available ligands (Hesselgesser et al. 1998; Meucci et al. 1998, 2000; Kaul and Lipton 1999; Kaul et al. 2007).

## 5 HIV-1-Induced Neurotoxicity

While progress is being made in unraveling the pathological processes, it remains a controversial issue how exactly HIV-1 infection provokes neuronal injury and death as well as neurocognitive and motor impairment (Kaul et al. 2005; Mattson et al. 2005; Gonzalez-Scarano and Martin-Garcia 2005; Kramer-Hammerle et al. 2005). There is general agreement that HIV does not infect postmitotic, mature neurons, but the mechanism of neuronal damage is in question.

Recent evidence suggests that viral clade may influence the occurrence of HIV-associated neuronal injury and development of HAND (reviewed in Gannon et al. (2011)). However, the HIV-1 genome of all clades encodes nine gene products which serve three major biological functions: (i) structure (Gag. polymerase (Pol), envelope protein (Env, gp120 and gp41)), (ii) regulation (transactivator of transcription (Tat), regulatory for expression of viral proteins (Rev)), or iii) accessory functions (viral protein u (Vpu), viral protein r (Vpr), viral infectivity factor (Vif), negative factor (Nef); recently reviewed in Ellis et al. (2007)). Most of these viral factors have been linked to distinct functions during infection and in the viral life cycle. Thus, whereas various HIV proteins affect the course of viral infection and disease, the envelope gp120 seems to play a major role in the induction of apoptosis in infected and uninfected bystander lymphocytes (Perfettini et al. 2005a, b). On the other hand, six HIV proteins have been reported to directly or indirectly affect neurons and glia (Ellis et al. 2007), and there exists ample evidence for neuronal injury by the five viral proteins Tat, Nef, Vpr, and the Env proteins gp120 and gp41 (Brenneman et al. 1988; New et al. 1997; Adamson et al. 1996; Piller et al. 1998; Koedel et al. 1999; Kaul et al. 2001, 2005; Mattson et al. 2005).

Human macrophages (and lymphocytes) provide suitable models to investigate HIV infection and potential neurotoxic effects in vitro and also in vivo, when injected into immunocompromised mice. However, using the intact virus and infection does not permit to study the contribution of a particular virus component to CNS injury. In order to study the injurious potential of HIV components in the brain in vivo, a number of transgenic mouse models have been generated (Toggas and Mucke 1996; Maung et al. 2012; Power et al. 2012; Gorantla et al. 2012). Transgenic mice expressing HIV/gp120 or Tat in their brains manifested several neuropathological features observed in AIDS brains, such as decreased synaptic and dendritic density, increased numbers of activated microglia, and pronounced astrocytosis (Toggas et al. 1994; Kim et al. 2003; Fitting et al. 2010). HIV/gp120 transgenic mice also developed significant physiological and behavioral changes, such as reduced escape latency, swimming velocity, and spatial retention at 12 months of age (Mucke et al. 1995; Krucker et al. 1998; Asensio et al. 2001; D'hooge et al. 1999). Following induction of Tat expression with doxycycline (Dox), inducible HIV-/Tat-transgenic (iTat-tg) mice displayed failure to thrive, impaired movement, and compromised spatial learning and memory (Kim et al. 2003; Carey et al. 2012; Fitting et al. 2012). HIV/Vpr-transgenic mice showed injury and loss of neurons and astrocytes but surprisingly lacked overt signs of neuroinflammation (Jones et al. 2007;

Power et al. 2012). Nevertheless, Vpr-tg mice developed learning and memory impairment. The relevance of the transgenic models was supported by findings obtained in two mouse models using HIV-infected human immune cells. In the first model, HIV-infected human macrophages were intracerebrally administered into mice with severe combined immunodeficiency (SCID) (Tyor et al. 1993; Persidsky et al. 1996), and the second model employed mice with a humanized hematopoietic system (Gorantla et al. 2010; Dash et al. 2011). The chimeric mice in the second model were infected with HIV-1 and followed for 15 weeks. The neuropathology in these latter two models using intact HIV-1 also resembled numerous pathological features of human AIDS brains and was in particular comparable to the brain injury observed in both HIV/gp120tg and HIV/iTat-tg mice.

Over time investigations using in vitro and in vivo models have generated evidence for at least two different hypotheses on how HIV-1 initiates neuronal damage in the brain. The hypotheses can be described as the "direct injury" hypothesis and the "indirect" or "bystander effect" hypothesis. These two hypotheses are by no means mutually exclusive, and while the available data support a role for both, an indirect form of neurotoxicity may predominate in a setting where monocytic, glial, and neuronal cells are present (Fig. 2) (Gartner 2000; Kaul et al. 2001, 2005; Mattson et al. 2005; Gonzalez-Scarano and Martin-Garcia 2005).

# 5.1 Direct Neuronal Toxicity

The hypothesis that HIV proteins can directly injure neurons without any contribution of nonneuronal cells (microglia and/or astrocytes) is supported by experiments showing that several viral proteins, including envelope protein gp120, are toxic in serum-free primary neuronal cultures (Meucci et al. 1998, 2000) and in neuroblastoma cell lines (Hesselgesser et al. 1998). The impact of neurotoxic factors secreted by non-neuronal cells is minimized in these experimental paradigms because serum-free neuronal cultures contain few if any nonneuronal cells and neuroblastoma or other neuron-type cell lines do not contain cells of other phenotypes.

In case of HIV envelope protein gp120, which interacts with several members of the chemokine receptor family (see above), the direct form of HIV-induced neuronal injury may be mediated by chemokine receptor signaling in the absence of CD4 (Hesselgesser et al. 1998). Indeed, experiments employing an inhibitor of CXCR4 can in some cases prevent HIV/gp120-induced neuronal apoptosis (Meucci et al. 1998; Kaul and Lipton 1999; Zheng et al. 1999; Meucci et al. 2000). In a different study, gp120 was found to interact at nanomolar concentrations with the glycine binding site of the *N*-methyl-*D*-aspartate-type glutamate receptor (NMDAR) (Fontana et al. 1997), suggesting an alternative mechanism by which HIV/gp120 may directly interfere with neuronal function and viability.

Interestingly, gp120 of the CXCR4-preferring HIV-1 strain IIIB can be endocytosed by neuronal axons and subsequently disturb axonal transport

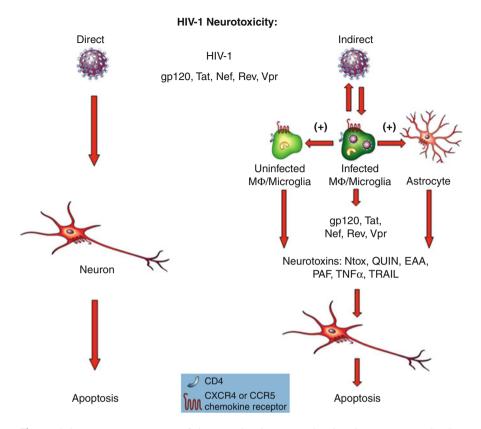


Fig. 2 Schematic representation of direct and indirect HIV-1-induced neurotoxicity. Studies using in vitro and in vivo models have generated evidence for at least two different hypotheses on how HIV-1 initiates neuronal damage in the brain: the "direct injury" hypothesis and the "indirect" or "bystander effect" hypothesis. These two hypotheses are by no means mutually exclusive. The "direct injury" hypothesis proposes that HIV and its proteins can directly damage and kill neurons without any contribution of nonneuronal cells, in particular microglia and/or astrocytes. The "indirect" or "bystander effect" hypothesis suggests that activation of macrophages and microglia by HIV infection or viral components and subsequent release of neurotoxins, such as excitotoxins and/or proinflammatory cytokines and mediators, provide the predominant trigger for neuronal injury and death. Macrophages ( $M\Phi$ ) and microglia are the major productively HIV-infected cell type in the brain and as such produce and release the virus and its components as well as cytokines and other factors that can stimulate uninfected MD/microglia and astrocytes. It is important to note that toxic viral proteins among factors released from microglia and excitatory glutamate and cytokines set free by astrocytes may all act in concert to provoke neurodegeneration and HAND, even in the absence of extensive viral invasion of the brain. Final neuronal loss seems to occur through apoptosis

(Mocchetti et al. 2012). Since neurotrophic support is critically dependent on this mode of transport, gp120 may not only compromise the endocytotic compartment but also neurotrophin function, and thus promote neuronal injury and demise. A follow-up study by the same group provided evidence that gp120 in addition

alters the neuronal processing of brain-derived neurotrophic factor (BDNF) in a fashion that facilitates neuronal deterioration (Bachis et al. 2012).

The viral regulatory protein Tat can freely diffuse through membranes and therefore interact with cellular binding partners on the surface as well as inside a cell (Fawell et al. 1994). Moreover, Tat can in principle interact with all cells although the consequences may vary between the individual cell types and depending on what Tat-interacting cellular factors are present. Tat can be taken up into neuron-like PC12 cells by a receptor-mediated mechanism (Liu et al. 2000) and may also have a direct effect on primary neurons by potentiating the response to excitotoxic stimuli (reviewed in (Mattson et al. 2005)). The various cell surface receptors that can interact with Tat include chemokine receptors, such as the CCR2, CCR3, and the HIV coreceptor CXCR4 (CD184), the low-density lipoprotein receptor-related protein (LRP), group I metabotropic glutamate receptors (mGluRs), and NMDA-type ionotropic glutamate receptors (Albini et al. 1998; Ghezzi et al. 2000; Xiao et al. 2000; Eugenin et al. 2007; Neri et al. 2007; Li et al. 2008). Inside a cell, Tat can affect glucagon synthase kinase (GSK) 3β and various transcription factors, such as nuclear receptor chicken ovalbumin upstream promoter transcription factor (COUP-TF) and AFF4 (Maggirwar et al. 1999; Rohr et al. 2000; He et al. 2010). A more recent study linked for the first time Tat to dysregulation of neuronal microRNAs (miR), including miR-128. The upregulation of miR-128 inhibited expression of the presynaptic protein SNAP25 (Eletto et al. 2008) and indicated a new way of how Tat may compromise synaptic function. Another report described a Tat-induced pathway to neurite retraction that required p73 and p53 and also led to accumulation of the pro-apoptotic Bcl-2 family protein Bax (Mukerjee et al. 2008). Finally, another direct pathway to neuronal injury initiated by Tat seems to involve activation of neuronal ryanodine receptors (RyR) by Tat, which causes a rapid loss of calcium  $(Ca^{2+})$  from the endoplasmic reticulum (ER) and induces the unfolded protein response (UPR) and mitochondrial hyperpolarization (Norman et al. 2008). The latter is associated with a compromised capability to buffer intracellular Ca<sup>2+</sup> and thus may affect neuronal excitability and synaptic activity.

Experiments using cultured hippocampal neurons suggested that the HIV protein Vpr may cause direct neurotoxicity through formation of a cation-permeable channel (Piller et al. 1998; Huang et al. 2000). Other studies found that HIV-1 Vpr induced apoptosis via a direct interaction with the mitochondrial adenine nucleotide translocator (ANT) and an effect on the permeability transition pore complex (PTPC) (Jacotot et al. 2000). However, the compromise of mitochondrial membrane potential and nuclear apoptosis triggered by Vpr were blocked in the presence of Bcl-2 (Jacotot et al. 2001). In human NT2 cell-derived neurons, Vpr was found to activate caspase-8 and trigger apoptosis (Pomerantz 2004). In a separate study, Vpr caused neuronal apoptosis accompanied by cytochrome c release, p53 induction, activation of caspase-9, and a diminishing effect on whole-cell currents. The same study found in Vpr-transgenic (tg) animals that the viral protein affected  $\gamma$ -aminobutyric acid (GABA)-ergic and cholinergic neurons in the basal ganglia besides a decreased immunoreactivity for subcortical synaptophysin and an

increase in activated caspase-3 (Jones et al. 2007). Of note, the in vivo neuropathology caused by Vpr in the tg mouse model may have resulted from direct and indirect effects (Power et al. 2012). Vpr was also reported to affect neuronal development by inducing mitochondrial dysfunction, reduced intracellular ATP concentration, compromised axonal transport of mitochondria, and consequent inhibition of axonal outgrowth (Kitayama et al. 2008).

HIV/Nef was also found to compromise neuronal survival, but the mechanism of the reported direct effect remained unknown (Trillo-Pazos et al. 2000). Of note, other studies found that Nef was present in astrocytes of HAD patients and induced apoptosis in astrocytes and neurons (He et al. 1997b). However, Nef triggered neuronal death indirectly because neurotoxicity required the presence of astrocytes and depended on their release of the CXCR3 ligand CXCL10/IP10 (van Marle et al. 2004).

A study investigating the potential neurotoxicity of HIV/Rev in vivo observed a lethal effect in mice that was linked to possible disturbance of cell membranes (Mabrouk et al. 1991). However, an in vitro study using nonneuronal cells found that Rev disturbed the structure of nucleoli and slowed the cell cycle during mitosis (Miyazaki et al. 1995). Thus, one may wonder whether Rev can affect besides HIV-infected, activated lymphocytes producing virus, also proliferating neural precursor cells.

While the absence of nonneuronal cells permits the investigation of potential direct effects of viral proteins on neurons, one ought to bear in mind that because of the absence of nonneuronal cells, a predominantly indirect effect cannot be detected. Notably, in the pathophysiological setting of the brain, neurons always encounter the potential toxins in the presence of glial and sometimes immune cells. Therefore, direct interaction of viral proteins with neurons, such as gp120 with neuronal chemokine receptors (Hesselgesser et al. 1997, 1998), may contribute to neuronal injury, but numerous additional studies suggest that activation of macrophage HIV coreceptors and subsequent release of neurotoxins, such as excitotoxins, chemokines, and/or proinflammatory cytokines, provide the predominant trigger for neuronal injury and death (Giulian et al. 1993; Kaul and Lipton 1999; Porcheray et al. 2006; O'Donnell et al. 2006; Cheung et al. 2008; Eggert et al. 2010; Medders et al. 2010).

#### 5.2 Indirect Neuronal Toxicity

Intact HIV-1, as well as picomolar concentrations of isolated viral envelope gp120, can induce neuronal death via CXCR4 and CCR5 receptors in cerebrocortical and hippocampal neurons and neuronal cell lines from humans and rodents (Hesselgesser et al. 1998; Iskander et al. 2004; Walsh et al. 2004; Meucci et al. 1998; Ohagen et al. 1999; Kaul and Lipton 1999; Chen et al. 2002; Garden et al. 2004; Kaul et al. 2007). In mixed neuronal/glial cerebrocortical or hippocampal cell cultures which model the cellular composition of brain regions affected by HIV, virus and gp120-induced apoptotic death appears to be mediated predominantly via the release of toxins from microglia and macrophages rather than by direct neuronal

damage (Brenneman et al. 1988; Iskander et al. 2004; Walsh et al. 2004; Giulian et al. 1990; Kaul and Lipton 1999; Chen et al. 2002; Garden et al. 2004; Medders et al. 2010). In fact, both the absence or inactivation of macrophages and microglia basically abrogates the neurotoxicity of HIV-1 and gp120 in mixed neuronal/glial cultures (Giulian et al. 1990; Giulian et al. 1993; Kaul and Lipton 1999; Chen et al. 2002; Garden et al. 2010). Since at least in vitro inhibition of microglial activation suffices to prevent neuronal death after gp120 exposure, it seems likely that stimulation of CXCR4 or CCR5 in macrophages and microglia is a prerequisite for the neurotoxicity of HIV-1 and gp120 (Ohagen et al. 1999; Kaul and Lipton 1999). This hypothesis is further supported by our observation that introduction of HIV coreceptor-expressing macrophages sufficed to restore neurotoxicity of gp120 in cerebrocortical cultures genetically deficient in both CXCR4 and CCR5 (Medders et al. 2010).

Chemokine receptors, including HIV coreceptors, are highly conserved between species (Maung et al. 2012). Therefore, our group recently studied the role of chemokine receptors in the neurotoxicity of gp120 using mixed neuronal/glial cerebrocortical cell cultures from rat and mouse. We observed that gp120 from CXCR4 (X4)-preferring as well as CCR5 (R5)-preferring and dual tropic HIV-1 strains all were able to trigger neuronal death, thus suggesting that both HIV coreceptors can mediate neurotoxicity, each alone and in combination (Kaul et al. 2007).

Experiments with cerebrocortical cell cultures from chemokine receptor knockout and wild-type mice addressed the role of each HIV coreceptor separately. Neurotoxicity of X4-preferring gp120 was strongly reduced or absent in CXCR4deficient cerebrocortical cultures, while gp120 from a R5-preferring HIV-1 strain showed no longer neurotoxicity in the absence of CCR5. Surprisingly, however, gp120 of the dual tropic HIV-1<sub>SF2</sub> exerted even greater neurotoxicity in CCR5 knockout cultures compared to wild-type or CXCR4-deficient cerebrocortical cells. Nevertheless, gp120 of HIV-1<sub>SF2</sub>, like all other tested gp120s, was no longer toxic to double KO cerebrocortical cultures lacking both chemokine receptors (Kaul et al. 2007). The finding that the CCR5 or CXCR4 knockout significantly decreased neurotoxicity by CCR5- or CXCR4-preferring gp120s, respectively, resembles the specificity of HIV coreceptors in human cells and thus confirmed the rodent system as a suitable model.

In addition, we observed earlier that the CCR5 ligands CCL4/MIP-1 $\beta$  and CCL5/RANTES protected neurons against gp120-induced toxicity (Kaul and Lipton 1999). Interestingly, this protective effect was strictly dependent on CCR5 but occurred no matter if gp120 preferred CXCR4 or CCR5 or both coreceptors. While the  $\beta$ -chemokines may have prevented CCR5-preferring gp120 from interacting with its receptor by competitive displacement, the blockade of CXCR4-mediated toxicity apparently involved heterologous cross-desensitization of this receptor via stimulation of CCR5 (Kaul et al. 2007). These findings were consistent with a primarily neurotoxic activation of CXCR4 by gp120. In contrast, CCR5 may at least in part have stimulated cytoprotective signals depending on the HIV-1 strain from which a given envelope protein originated and depending on

whether or not antiviral and cytoprotective  $\beta$ -chemokines were present (Cocchi et al. 1995; Kaul et al. 2007).

Interestingly, in macrophages, HIV-1 infection and exposure to just the envelope protein gp120 seem to trigger a similar neurotoxic phenotype (Giulian et al. 1990, 1993). Tat and Nef each alone also causes activation of mononuclear phagocytes and release of cytokines as it can occur upon viral infection (Federico et al. 2001; Olivetta et al. 2003; Sui et al. 2006). In contrast, Vpr seems to promote a rather anti- than proinflammatory phenotype despite evidence that this viral protein plays an important role in HIV production in macrophages (Herbein et al. 2010; Power et al. 2012).

In any case, uninfected macrophages and microglia can be stimulated by factors released from infected cells. These factors include cytokines, such as IL-1 $\beta$ , IL-6, and TNF $\alpha$ , but also TNF-related apoptosis-inducing ligand (TRAIL), matrix metalloproteinases (MMPs), arachidonic acid metabolites, and platelet-activating factor (PAF), and shed viral proteins, such as gp120, Tat, Vpr, and Nef (Wahl et al. 1989; Achim et al. 1993; Gelbard et al. 1994; Miura et al. 2003a, b; Ryan et al. 2004; Conant et al. 1998; Johnston et al. 2000; Herbein et al. 2010). Variations of the HIV-1 gp120, in particular in its V1, V2, and V3 loop sequences, have been implicated in modulating the neurotoxicity of macrophages and microglia (Power et al. 1994, 1998). Activated microglia affect all cell types in the CNS, resulting in further upregulation of cytokines, chemokines, and endothelial adhesion molecules within the brain (Gartner 2000; Kaul et al. 2001, 2005; Gonzalez-Scarano and Martin-Garcia 2005; Kramer-Hammerle et al. 2005).

Some of the secreted factors from activated microglia or macrophages and astrocytes may directly or indirectly contribute to neuronal damage and apoptosis as discussed above for viral proteins. Directly neurotoxic host factors released from HIV-infected or activated microglia and macrophages include excitatory amino acids (EAAs) and related substances, such as quinolinate, a not completely characterized amine compound named "Ntox" but potentially also TRAIL (Giulian et al. 1990; Kramer-Hammerle et al. 2005; Lipton et al. 1991; Heyes et al. 1991; Kaul et al. 2001, 2005; Gonzalez-Scarano and Martin-Garcia 2005; Yeh et al. 2000; Miura et al. 2003a, b; Ryan et al. 2004). EAAs and quinolinic acid can trigger neuronal apoptosis through a process known as excitotoxicity. This detrimental process involves excessive Ca<sup>2+</sup> influx and free radical (nitric oxide and superoxide anion) formation by overstimulation of glutamate receptors (Lipton et al. 1991), activation of stress-associated protein kinases and caspases, and production of proinflammatory lipids (Kaul et al. 2001; Mattson et al. 2005; Gonzalez-Scarano and Martin-Garcia 2005; Kramer-Hammerle et al. 2005; Jones and Power 2006; Nicotera et al. 1997).

It is important to note that toxic viral proteins among factors released from microglia and glutamate and cytokines set free by astrocytes may all act in concert to provoke neurodegeneration and HAND, even in the absence of extensive viral invasion of the brain. Furthermore, HAND and HAD might share the critical involvement of neuroinflammation and microglial activation with several other neurodegenerative diseases, such as Alzheimer's disease, multiple sclerosis, Parkinson's disease, and even frontotemporal lobe dementia (Block and Hong 2005).

# 5.3 Downstream Cellular and Molecular Mechanisms in HIV-1 Neurotoxicity

Beyond infection by intact HIV-1, six of the viral protein components directly or indirectly affect immune cells, neurons, and glia (Ellis et al. 2007), and ample evidence exists for neuronal injury by the five viral proteins Tat, Nef, Vpr, and the Env proteins gp120 and gp41 (Brenneman et al. 1988; New et al. 1997; Adamson et al. 1996; Piller et al. 1998; Koedel et al. 1999; Kaul et al. 2001, 2005; Mattson et al. 2005). Therefore, it is not surprising that the available information suggests that HIV can trigger a plethora of cellular responses and signaling mechanisms. However, it appears that the various viral components that exert cyto- or neurotoxicity do so, at least in part, by converging mechanisms even if each component may start out its contribution to pathogenesis by interacting with different host cell factors. Currently recognized converging mechanisms include neuroinflammation, oxidative stress, interference with the cell cycle machinery, excitotoxicity, and apoptosis.

Recently, we and others found that p38 mitogen-activated protein kinase (MAPK) plays an essential role in the induction of neuronal injury and apoptotic death by HIV-1 gp120, Tat, and the intact virus (Kaul and Lipton 1999; Kaul et al. 2007; Choi et al. 2007; Medders et al. 2010; Singh et al. 2005; Sui et al. 2006; Eggert et al. 2010). Activation of microglia and macrophages by HIVgp120 via CD4, chemokine coreceptors, and downstream p38 MAPK is crucial to neurotoxin production (Kaul and Lipton 1999; Medders et al. 2010). Similarly, others reported that HIV-1 infection as well as Tat and gp120 all activate p38 MAPK and upstream, the mixed lineage kinase (MLK) 3 in infected or stimulated, neurotoxic monocytes and macrophages (Sui et al. 2006; Eggert et al. 2010). Moreover, we have observed that p38 MAPK signaling is essential upon exposure to HIV gp120 for both the neurotoxic phenotype of macrophages and microglia and the subsequent induction of neuronal apoptosis by macrophage toxins (Medders et al. 2010). The requirement of p38 MAPK activity in both macrophages and neurons for the occurrence of HIV gp120 neurotoxicity therefore parallels the indispensability previously reported for the transcription factor p53 (Garden et al. 2004).

Increased activity of p38 MAPK has been linked to inflammation of the brain, such as that seen in HIVE patients and Alzheimer's disease (AD) (Kaminska et al. 2009). Furthermore, neurons showed increased activity of p38 MAPK when they were cocultured with LPS-stimulated microglia, and this kinase signaling was essential to induction of neuronal death (Xie et al. 2004). Amyloid- $\beta$ -induced activation of p38 MAPK resulted in upregulation of proinflammatory cytokines TNF $\alpha$ , immunologic, inducible nitric oxide synthase (iNOS), and interleukin (IL)-1 $\beta$  and subsequent neuronal death (Delgado et al. 2008). Our and other groups showed that inhibition of p38 MAPK activity in vitro with pharmacological

inhibitor SB203580 or dominant-negative  $p38\alpha$ AF (TGY substituted with AGF) prevented neuronal death induced by HIV-1 gp120 or glutamate (Kaul and Lipton 1999; Singh et al. 2005; Kaul et al. 2007; Choi et al. 2007; Chaparro-Huerta et al. 2008). HIV-1 Tat and gp120 both were found to activate MLK 3, which controls p38 MAPK and JNK pathways, in neurons (Sui et al. 2006; Gallo and Johnson 2002). Inhibition or MLK 3 protected neurons from toxicity of gp120, Tat, and HIV-1-infected macrophages (Sui et al. 2006; Eggert et al. 2010). Of note, while pharmacological inhibition of p38 MAPK or JNK can both independently ameliorate the toxicity of Tat in cerebellar granular neurons, only blockade of p38 MAPK, but not JNK, could prevent Tat-induced death in striatal neurons (Singh et al. 2005; Sui et al. 2006).

In neurons, p38 $\alpha$  MAPK activity is regulated by the intracellular Ca<sup>2+</sup> concentration and the small GTPase Rho. The concentration of intracellular Ca<sup>2+</sup> in turn is linked to the activity of the NMDAR-regulated ion channel and excitotoxicity (Semenova et al. 2007). A number of studies have suggested that excitotoxicity itself constitutes one important mechanism of HIV-1-induced neurotoxicity (Dreyer et al. 1990; O'Donnell et al. 2006; Kaul et al. 2001; Lindl et al. 2010). The link between glutamate receptors, influx of Ca<sup>2+</sup>, Rho, and activation of p38 $\alpha$  certainly suggests that in neurons synaptic activity may be a major regulator of the kinase's baseline activity. Our studies indicate that inhibition of p38 MAPK prevents neuronal death in the context of HIV/gp120-induced, macrophage-mediated neurotoxicity, but surprisingly, neurons surviving in the absence of kinase inhibition display a higher level of active p38 MAPK compared to control cells not exposed to neurotoxins (Medders et al. 2010).

Macrophages and microglia infected with HIV-1 or stimulated by viral gp120, Tat, or Nef release neurotoxins, some of which stimulate the NMDAR, an ionotropic glutamate and neurotransmitter receptor (Dreyer et al. 1990; Chen et al. 2002; Smith et al. 2001; Sui et al. 2007). NMDAR antagonists can ameliorate neuronal cell death in vitro due to HIV-infected macrophages or purified recombinant gp120 (Dreyer et al. 1990; Chen et al. 2002) and in vivo in gp120 transgenic mice (Toggas et al. 1996). Under physiological conditions, activation of ionotropic glutamate receptors in neurons initiates a transient depolarization and excitation. However, extended or excessive NMDAR activation results in excitotoxicity because the prolonged channel opening leads to a sustained elevation of the intracellular Ca2+ concentration and thus compromises mitochondrial function and cellular energy metabolism, which in turn causes the generation of free radicals (Lipton and Rosenberg 1994; Doble 1999; Olney 1969). In case the initial excitotoxic insult is fulminant, for example, in the ischemic core of a stroke, the neurons die early from loss of ionic homeostasis, resulting in acute swelling and cell lysis (necrosis). If the insult is milder, as seen in several neurodegenerative disorders, including HAND/HAD, neurons may enter a delayed death pathway known as programmed cell death or apoptosis (Nicotera et al. 1997; Petito and Roberts 1995). Neuronal apoptosis following exposure to gp120-induced macrophage toxins or after a direct excitotoxic insult involves Ca<sup>2+</sup> overload, activation of p38 MAPK and p53, release of cytochrome c and other molecules such as apoptosis-inducing factor (AIF) from mitochondria, activation of caspases, free radical formation, lipid peroxidation, and chromatin condensation (Tenneti et al. 1998; Kaul and Lipton 1999; Asensio and Campbell 1999; Garden et al. 2002, 2004; Kaul et al. 2007). The increase of intracellular  $Ca^{2+}$  has also been linked to the detrimental activation of the  $Ca^{2+}$ -dependent protease Calpain and the activating, proteolytic cleavage of the cell cycle-dependent kinase (Cdk)-5 (Wang et al. 2007). Excessive activation of  $Ca^{2+}$ -dependent proteolysis has also been implicated in synaptic injury where the scaffolding inside of pre- and postsynaptic endings may be compromised and thus disturb timely trafficking and localization of vesicles as well as receptor signaling (Lindl et al. 2010). In line with  $Ca^{2+}$ -dependent oxidative stress is a different study reporting that in human neurons, CXCR4 mediates the toxic effect of gp120 via a process involving the synthesis of ceramide and NADPH-dependent production of superoxide radicals (Jana and Pahan 2004).

Oxidative stress is a well-recognized cellular response triggered by infections. inflammation, as well as excitotoxicity and occurs in the context for HIV infection or exposure to viral proteins (Roederer et al. 1992; Walmsley et al. 1997; Shi et al. 1998; Corasaniti et al. 2000; Mollace et al. 2001; Boven et al. 1999; Haughev et al. 2004; Louboutin et al. 2007; Agrawal et al. 2009). Oxidative damage occurs in proteins, lipids, and nuclear DNA and can therefore contribute in several ways to neuronal deterioration (Louboutin et al. 2010; Haughey et al. 2004; Bonfoco et al. 1995; Zhang et al. 2012). Interestingly, gp41 in infected cells was found to correlate with the expression of iNOS in AIDS brains and cause neurotoxicity in vitro in an NO-dependent fashion (Adamson et al. 1996). Oxidative stress alters proteins by causing carbonylation and affects lipid metabolism. The latter leads to generation of distinct patterns of ceramide, sphingomyelin, and hydroxynonenal (HNE) detectable in brain tissue and CSF of HAND patients (Turchan et al. 2003; Haughey et al. 2004; Mattson et al. 2005). In line with oxidative stress is the upregulation of the transcription factor ATF-6ß and of the ER chaperone BiP indicating ER stress in neurons and astrocytes of brains from neurocognitively impaired HIV patients (Lindl et al. 2007). In a more recent follow-up study, the same group linked ATF-6ß and BiP to an "integrated stress response," which brains of HAND/HAD patients may share with other neurodegenerative diseases (Akay et al. 2012).

Autophagy allows cells to remove exhausted or damaged organelles or to replenish their resources under starvation conditions and appears to be an essential mechanism to maintain cellular homeostasis. Recent studies strongly suggest that autophagy is disturbed in HIV infection and neurodegenerative diseases, including HIVE and HAND (Espert et al. 2006; Alirezaei et al. 2008; Zhou et al. 2011; Hui et al. 2012). Another mechanistic link to HIV neurotoxicity is suggested by the observation that both TNF $\alpha$  and excitotoxic glutamate concentrations seem to compromise neuronal autophagy and in this way also promote cell death (Alirezaei et al. 2008).

HIV-infected or gp120-activated microglia also release inflammatory cytokines, such as TNF $\alpha$  (Achim et al. 1993; Wesselingh et al. 1997; Persidsky et al. 1997). TNF $\alpha$  directly affects glutamate neurotransmission in a p38 MAPK-dependent

manner by increasing the synaptic expression of certain glutamate receptors (a-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid receptor; AMPAR) and inhibiting long-term potentiation (LTP) (Butler et al. 2004; Pickering et al. 2005). Interestingly, a significant change in the expression at the RNA level of a number of ion channels was reported for brains of HIV patients with various degrees of HAND in comparison to uninfected controls (Gelman et al. 2004). Those disturbances of the ion channel repertoire were likely to affect excitability of neuronal membranes, neurotransmission, as well as the propagation of action potentials in neurons. Therefore, it was proposed that a channelopathy could be underlying the development of HAND and its most severe form, HAD. In any case,  $TNF\alpha$  can inhibit astroglial glutamate transporters and thus promote neurotoxicity by facilitating glutamate excitotoxicity (Pickering et al. 2005) or even provoke neurotransmitter release from glial cells (Benveniste and Benos 1995; Bezzi et al. 2001). Interestingly, TNF $\alpha$  can synergize with HIV/Tat to initiate neuronal death, but antioxidants block the process (Ohagen et al. 1999). Finally, it remains possible that TNFa activates caspases in neurons via TNF $\alpha$  receptor-1 (TNFR1). TNFR1 is expressed on at least some neurons, and it can initiate caspase-8 activation, which in turn may activate caspase-3, leading to apoptosis. We found that neutralization of TNF $\alpha$  with antibody or pharmacological inhibition of caspase-8 prevented the neurotoxicity of HIV/gp120 in cultured cerebrocortical neurons (Garden et al. 2002).

Supporting a prominent role of neuronal apoptosis in the development of HAND and HAD, activated caspase-3 and p53 were prominently detected in neurons of brains from neurocognitively impaired HIV patients (Garden et al. 2002). Furthermore, in vitro, neuronal caspase-3, caspase-8, and caspase-9 were found to be necessary for HIV-1/gp120 to cause neurotoxicity, and p53 was essential not only in neurons but also in microglia (Garden et al. 2002, 2004; Tun et al. 2007). Separate studies suggested that CXCR4 and p53 are connected through signaling pathways that mediate toxic or protective mechanisms depending on whether gp120 or SDF-1, the receptor's natural binding partner, serves as the ligand (Khan et al. 2005). Contrary to gp120 of HIV1IIIB, CXCL12/SDF-1 was neuroprotective in this experimental paradigm. The  $\alpha$ -chemokine activated Akt and MAPKs (Khan et al. 2004) and regulated the expression and localization of cell cycle proteins (Khan et al. 2003, 2005). CXCL12/SDF-1 increased acetylation of p53 and p21 as well as the expression of retinoblastoma protein (Rb) while reducing the amount of phosphorylated Rb in the nucleus. Together with a reduction of the activity of the transcription factor E2F1, an overall anti-apoptotic effect was observed. In contrast, envelope protein of HIV-1<sub>IIIB</sub> triggered activation of Apaf-1 and promoted cell death. Besides these in vitro findings, a role for a disturbed cell cycle machinery in the development of HAND was supported by abnormal changes in the expression pattern of the same cell cycle proteins in *postmortem* brains of nonhuman primates with SIV encephalitis and humans with HIVE (Jordan-Sciutto et al. 2002). Interestingly, these changes in cell cycle proteins also correlated with the presence of activated microglia and macrophages. Moreover, cell cycle inhibition has been shown to protect neurons in excitotoxic circumstances from induction of apoptosis (Park et al. 2000).

#### 6 Effects of HIV-1 on Neurogenesis

Compared to the discovery of HIV-induced neurotoxicity, the appreciation of the virus' effect on neurogenesis is a relatively recent development. Notably, HIV primarily seems to compromise the proliferation of neural progenitors but not kill the cells (Krathwohl and Kaiser 2004a, b; Okamoto et al. 2007; Peng et al. 2008). Therefore, HIV does not exert a lethal cytotoxic effect on neural precursors, but rather exacerbates the effects of the above-discussed toxicity in postmitotic and terminally differentiated neurons by impairing neurogenesis and brain homeostasis.

It is widely accepted that HIV-1 fails to productively infect neurons, but it has been reported that neural progenitor cells are permissive to the virus (Lawrence et al. 2004; Mattson et al. 2005; Schwartz and Major 2006). However, infection seems dispensable as HIV-1 gp120 and chemokines suffice to disturb human or rodent neural progenitor cells (Krathwohl and Kaiser 2004a, b; Okamoto et al. 2007). One study showed that chemokines promoted the quiescence and survival of human neural progenitor cells via stimulation of CXCR4 and CCR3 and a mechanism that involved downregulation of extracellularly regulated kinase-1 and kinase-2 (ERK-1/ERK-2) with simultaneous upregulation of the neuronal glycoprotein reelin (Krathwohl and Kaiser 2004a). Exposure to intact HIV-1 also caused quiescence of neural progenitors through engagement of CXCR4 and CCR3. In contrast to the chemokines, the viral envelope protein gp120 downregulated ERK-1/ERK-2 but had no effect on reelin (Krathwohl and Kaiser 2004b). Interestingly, the effects of both the chemokines and HIV-1/gp120 were reversible and could be inhibited with recombinant apolipoprotein E3 (ApoE3), but not ApoE4. Finally, fewer adult neural progenitor cells were detected in the dentate gyrus of the hippocampus in *postmortem* brain specimen from HAD patients than from non-demented and uninfected controls (Krathwohl and Kaiser 2004b).

Cultures of primary rodent and human neural progenitor cells obtained from fetal or adult tissue express CXCR4 and CCR5 besides the neural progenitor marker nestin and readily undergo cell division. After several rounds of proliferation, the progenitors exit the cell cycle and begin to express neuronal markers such as BIII-tubulin (TuJ1). We found that exposure to HIV-1/gp120 in vitro reduced the proliferation of adult progenitors without producing cell death. Others observed a virtually identical effect brought about by HIV Tat (Mishra et al. 2010). Similar to what others observed in brain specimen from HAD patients, we found a reduction of proliferating neural progenitors in the hippocampal dentate gyrus of transgenic mice that express HIV/gp120 in the brain in comparison to non-transgenic controls (Okamoto et al. 2007). Accounting for these effects, we found that gp120 inhibited proliferation of neural progenitor cells through activation of a pathway that involved p38MAPK, MAPK-activated protein kinase-2, a cell cycle checkpoint kinase, and Cdc25B/C which in turn caused a delay of the cell cycle in the G1 phase. The decrease in neural progenitor proliferation caused by gp120 constituted an impairment in neurogenesis because there were fewer progenitor cells available to differentiate into neurons (Okamoto et al. 2007). Neurogenesis in the dentate gyrus of the hippocampus has been implicated in learning and memory formation, neurocognitive functions which are compromised in HAND. The apparent ability of HIV-1 or its component gp120 and Tat to interfere with the normal function of neural progenitor cells suggested the possibility that HAND might develop as a consequence not only of injury and death of existing neurons but also due to virus-induced disturbance of brain homeostasis and renewal mechanisms.

## 7 Comorbidity Factors of HAND and HIV-1 Neurotoxicity

HIV infection, HAND, and AIDS are frequently associated with the use of recreational drugs, such as methamphetamine, cocaine, and opiates (Rippeth et al. 2004; Urbina and Jones 2004; Kapadia et al. 2005; Mitchell et al. 2006; Nath 2010; Beyrer et al. 2010; Gannon et al. 2011). Intravenous drug use not only is a risk factor for contracting HIV through needle sharing but also promotes coinfection with hepatitis C virus (HCV) (Gatignol et al. 2007; Clifford et al. 2009; Cherner et al. 2005). The scientific literature provides ample evidence that both abused drugs and infectious agents, such as HIV and HCV, compromise the immune and nervous system, but this still-expanding topic is beyond the scope of this chapter. However, there exist excellent reviews addressing the apparent intersection of HAND and HIV-induced neurotoxicity and impaired neurogenesis with drug abuse (Hauser et al. 2006; Ferris et al. 2008; Nath 2010).

HIV infection has become a chronic rather than acute disease due to efficient control of peripheral viral replication by cART and the treatment of opportunistic infections. With the improved survival times, the population of HIV patients has begun to age. While aging is a normal physiological process, evidence is accumulating that HIV infection may lead to earlier occurrence of aging symptoms physically, immunologically, and mentally (Deeks 2009; Brew et al. 2009). Thus, aging is becoming a comorbidity factor associated with long survival of HIV infection. In reverse, the virus can also be seen as a comorbidity factor of aging that accelerates an otherwise normal but later ensuing increase of frailty.

#### 8 Conclusion

HIV-1 infection continues to trigger neurological problems that can culminate in frank dementia, despite efficient virological control by cART. Currently, strong evidence suggests that HIV-1 can strike at the brain in at least two ways: (i) by causing toxicity leading to neuronal injury and death and (ii) by impairing neurogenesis without killing neural precursors. Both of the pathogenic mechanisms share the involvement of neuroinflammatory mediators, and both processes seem to be influenced by host and viral factors, including the interferon response, age, and viral subtypes, as well as treatment and comorbidity factors, such as drug abuse. Microbial translocation may enhance systemic and central inflammation during HIV-1 infection and thus promote the development of HAND. Thus, despite viral control and life-prolonging effects of cART, neuroinflammation and concomitant

activation of macrophages and microglia accompany the continued development of HAND in HIV patients. Besides intact HIV-1, six viral proteins have been reported to directly or indirectly affect neurons and glia and cause neurotoxicity: Tat, Nef, Vpr, Rev, and the Env proteins gp120 and gp41. Experimental evidence generated at least two different hypotheses on how HIV-1 initiates neuronal damage in the brain: (i) the "direct injury" hypothesis and (ii) the "indirect" or "bystander effect" hypothesis. These two hypotheses are by no means mutually exclusive. However, macrophages and microglia constitute the primary productively infected cell type in the brain, suggesting that an indirect form of neurotoxicity may predominate in a setting where monocytic, glial, and neuronal cells are present. Considering that beyond infection by intact HIV-1, six of the viral proteins trigger neuronal injury, it is not surprising that the available information suggests a plethora of cellular responses and signaling mechanisms. However, it appears that the various viral components that exert cyto- or neurotoxicity do so, at least in part, by converging mechanisms even if each component may start out its contribution to pathogenesis by interacting with different host cell factors. Currently recognized converging mechanisms in HIV neurotoxicity include neuroinflammation, MAPK signaling, altered gene expression, oxidative stress, interference with the cell cycle machinery, excitotoxicity, synaptic injury, and neuronal apoptosis.

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# **Cancer-Mediated Neurotoxicity**

## Andrzej Stepulak, Radosław Rola, and Krzysztof Polberg

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#### Abstract

The central nervous system is particularly susceptible to cancer, and its treatment side effects result in significant neurologic morbidity and mortality. Conversely, recent improvements in treatment protocols have extended average life expectancies in various subgroups of cancer patients. As a result of the extended survival, quality of survival is becoming much more important, and attention is increasingly being paid to the disease-free period between treatment and death. The majority of the research, however, focuses on potential side effects of various therapeutic modalities. The following contribution is devoted to

A. Stepulak (🖂)

Department of Otolaryngology, MSW Hospital, Lublin, Poland e-mail: andrzej.stepulak@gmail.com

R. Rola

Department of Neurological Surgery, Medical University of Lublin, Lublin, Poland

Department of Physiopathology, Institute of Agricultural Medicine, Lublin, Poland e-mail: rola.radoslaw@gmail.com

K. Polberg

Department of Biochemistry and Molecular Biology, Medical University of Lublin, Lublin, Poland

Department of Otolaryngology, MSW Hospital, Lublin, Poland e-mail: kpolberg@poczta.onet.pl

exploration of a relatively new concept of tumor-related neurotoxicity with special emphasis on the role of glutamate and glutamate-dependent pathways in tumor development and progression as well as in glutamate-dependent excitotoxicity and its consequences.

#### Keywords

Cancer • Glutamate • Neurotoxicity • Receptors

List of Abbreviations		
	AMPA	α-amino-3-hydroxy-5-methyl-4 isoxazolepropionate
	EAAT	Excitatory amino acid transporter
	ECM	Extracellular matrix
	FAK	Focal adhesion kinase
	GDH	Glutamate dehydrogenase
	Glu	Glutamate
	HAA	Hydroxyanthranilic acid
	HDI	Histone deacetylase inhibitor
	IDO	Indoleamine 2,3-dioxygenase
	iGluR	Inotropic glutamate receptor
	KA	Kainate
	KYNA	Kynurenic acid
	LTD	Long-term depression
	LTP	Long-term potentiation
	MAGUK	Membrane-associated guanylate kinase
	mGluR	Metabotropic glutamate receptor
	NMDA	<i>N</i> -methyl-D-aspartate
	NOS	Nitric oxide synthase
	PIC	Picolinic acid
	QA	Quinolinic acid
	TMZ	Temozolomide
	TSC	Tumor stem cell

## 1 Introduction

According to GLOBOCAN database of the International Agency for Research on Cancer, almost 12.7 million new cases of cancer were reported worldwide in 2008, resulting in more than 7.6 million deaths (Ferlay et al. 2010). A significant percentage of those patients are expected to develop complications in sites distant to the original tumor location. The nervous system is particularly susceptible to such complications that result in significant neurologic morbidity and mortality. These effects may be direct, with direct cancer involvement of the brain, spine, or peripheral nervous system (PNS) or indirect as in paraneoplastic neurologic syndromes (Giglio and Gilbert 2010) with symptoms ranging from delirium, amnesia,

aphasia, psychotic disturbance, sensory disturbance, ataxia, and autonomic dysfunction to eve movement disorders (Bataller and Dalmau 2004). Still, the impairments of cognitive functions such as language, memory, visuospatial perception, time-space orientation, and attention or executive functions are the most prevalent (Reid-Arndt et al. 2010). From this perspective a distinct subpopulation of cancer patients is patients with brain tumors, both primary and metastatic ones who, in fact, show signs of cognitive decline in almost all of the cases (Hottinger et al. 2009). As a matter of fact, a cognitive dysfunction often is present at diagnosis in many of those patients, more frequently in rapid-growing tumors such as glioblastoma than in slow-growing ones such as low-grade gliomas (Correa 2010). On the other hand, improvements in treatments have extended average life expectancies in various subgroups of patients with brain tumors, although in most instances the patient cannot be cured and will eventually die from the disease (DeAngelis 2001). As a result of the extended survival, quality of survival is becoming much more important, and attention is increasingly being paid to the disease-free period between treatment and death (Weitzner and Meyers 1997).

The exact pathophysiology of cognitive deficits in patients with brain tumors is not entirely understood. Studies have shown that tumor itself, its progression, tumor-related neurological complications, and epilepsy can cause them (Klein et al. 2002; Aoyama et al. 2007). Diaschisis, in which there is dysfunction of brain areas connected to the original lesion, owing to altered metabolism and blood flow, might also contribute (Taphoorn and Klein 2004). On top of that, disturbances in connectivity between functional networks in the brain have also been reported (Bartolomei et al. 2006). The question remains, however: what is the underlying mechanism responsible for these disturbances?

A recent discovery of glutamate as a neurotrophic factor for gliomas, acting in autocrine or paracrine fashion, not only revealed a potential signaling pathway important for glioma invasion and established a new tool for cancer treatment but also opened a new avenue of research into the mechanisms responsible for neuronal network impairments found in brain tumor patients. Glial tumors, especially malignant gliomas, release significant amounts of Glu into extracellular space, which acts as a growth factor for malignant cells. In the last decade evidence has emerged implicating a role for glutamate as a signal mediator that stimulates cancer cell proliferation and motility via activation of glutamate receptors (GluRs) (de Groot and Sontheimer 2011). Moreover, excess of glutamate causes neurodegeneration of the peritumoral tissue, enabling tumor progression (de Groot and Sontheimer 2011). On the other side, elevated glutamate concentrations in the brain exert neurotoxicity, resulting in neuronal excitotoxicity and cell death, enabling brain tumors to expand, and cause seizures in the tumor-bearing patients (Sontheimer 2008; Buckingham et al. 2011). Importantly, glutamate is considered as one of the major excitatory mediators in the central nervous system (CNS). It affects several physiological processes such us learning, memory, or behavior (Peng et al. 2011). Excessive activation of glutametergic system (high levels of Glu and activation of respective receptors) results in neural cells death called excitotoxicity that is involved in pathophysiology of various neurodegenerative diseases and other CNS pathologies (epilepsy, hypoxia, stroke) (Lau and Tymianski 2010). Thereby, gliomaderived glutamate seems to be important endogenous neurotoxin.

Glutamate activates two principal classes of receptors: ionotropic (iGluRs) and metabotropic (mGluRs) glutamate receptors. The mGluR receptor family comprises of eight subtypes (mGluR1-8) that are subdivided into three groups based on the sequence similarities and G-protein coupling to specific signal transduction mechanisms. mGluR1 and mGluR5, which are coupled to the phospholipase C, belong to group I. Group II consists of mGluR2 and mGluR3, whereas group III contains mGluR4, mGluR6, mGluR7, and mGluR8, which are all negatively coupled to adenylate cyclase. mGluRs function as dimers, with two glutamate molecules being required for full receptor activation (Shin et al. 2008; Kew and Kemp 2005). In neurons mGluR receptors modulate synaptic transmission and are involved in activity-dependent modification of synaptic transmission such as long-term potentiation and long-term depression, whereas in glial cells they exert a variety of effects that are crucial for glial function and glial-neuronal interaction under physiologic and pathologic conditions (D'Antoni et al. 2008). iGluRs, on the other hand, are subdivided into three groups according to structural similarities and synthetic agonists that activate them: N-methyl- $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionate **D**-aspartate (NMDA), (AMPA), and 2-carboxy-3-carboxymethyl-4-isopropenylpyrrolidine (kainate, KA) receptors.

*N*-methyl-D-aspartate (NMDA) tetrameric receptors consist of two obligatory NR1 subunits and two of four types of regulatory subunits: NR2A, B, C, and D. The NR1 subunit is necessary for calcium conductivity of the channel, while the NR2 and NR3 subunits determine electrophysiological and pharmacological properties of the receptor. Activation of NMDA receptor requires binding of two agonists: glutamate and glycine. Concurrent membrane depolarization abrogates the magnesium block, which allows calcium and sodium cations to enter the cell. NR3 subunits may assemble with NR1 subunits in order to create a functional glycine receptor that is not affected by glutamate. NMDA receptors are implicated in synaptic transmission and neuronal plasticity phenomena such as long-term potentiation (LTP) and long-term depression (LTD) (Kew and Kemp 2005; Traynelis et al. 2010).

AMPA receptors are homo- or heterotetramers composed of subunits known as GluR1–GluR4. They mediate fast excitatory synaptic signaling in the brain and are involved in activity-dependent modulation of synaptic plasticity. AMPA receptors gate Na+/Ca<sup>2+</sup> in response to ligand binding. Conductance and kinetic properties of the channel depend upon subunit composition. Influx of ions causes a fast excitatory postsynaptic response in neurons, and Ca<sup>2+</sup> can also activate second messenger pathways. The GluR2 subunit plays a critical role in the determination of the permeability of AMPARs. The genomic DNA of the GluR2 subunit contains a glutamine (Q) residue at amino acid 607, but during the process of nuclear RNA editing, glutamine is replaced by arginine (R) in the vast majority of neuronal cells, which results in a very low calcium permeability for receptors containing the

GluR2 subunit (Madden 2002; Kew and Kemp 2005). The kinetics and amplitude of the excitatory synaptic response are determined by the biophysical properties of the receptor subunit combination (Traynelis et al. 2010).

KA receptors are composed of tetrameric assemblies of GluR5–7 and KA1/2 subunits. GluR5–7 subunits can form homomeric functional receptors as well as combine with KA1 and KA2 to form heteromeric receptors with distinct pharma-cological properties. KA1 and KA2 subunits do not form homomeric functional receptors. Similar to GluR2, GluR5 and GluR6 undergo glutamine/arginine editing (Contractor et al. 2011).

The glutamate cellular system includes also a number of proteins transporting glutamate, namely, high-affinity sodium and potassium-coupled excitatory amino acid transporters (EAAT1/GLAST, EAAT2/GLT-1, EAAT3/EAAC1, EAAT4, EAAT5). The major function of these transporters is to maintain the extracellular glutamate concentration at a low, nontoxic level in order to prevent overstimulation of glutamate receptors. Glu uptake occurs primarily through EAAT1 and EAAT2 transporters present on membranes of nonmalignant astrocytes (de Groot and Sontheimer 2011).

#### 2 Developments in the Field

Neurotoxic effects of glutamate and its relevance for the CNS as a potential neurotransmitter were recognized for the first time over 60 years ago. In 1952, severe convulsions were observed following glutamate administration into the central nervous system (CNS) of dogs and monkeys (Hayashi 1952), and in 1957 the neurotoxic effects of sodium-L-glutamate on the inner layers of the retina were described (Lucas and Newhouse 1957). In 1959, direct evidence that Glu depolarized individual neurons of the spinal cord was provided (Curtis et al. 1959). With the identification of natural or synthetic substances (NMDA, AMPA, KA) capable to discriminate and activate different glutamate receptors, followed by receptor antagonist/agonist development and mapping of the various iGluR subtypes to distinct CNS regions by radioligand binding techniques, and finally with molecular receptor characterization based on structure-function analysis, significant progress was made in elucidating the role of GluRs in physiology and CNS disorders (Bowie 2008).

Various studies have proven glutamate to be the major excitatory neurotransmitter in the brain that plays a key role in learning and memory, thanks to its role in the induction and maintenance of long-term potentiation (LTP) and long-term depression (LTD). For that, its signaling pathways have been extensively studied in the CNS. Accordingly, aside from the physiological functions of Glu, much effort has been made in order to resolve its mechanism of action causing neurotoxicity, termed excitotoxicity (Lau and Tymianski 2010).

Excessive glutamate levels cause neuronal dysfunction and degeneration by the complex excitotoxic mechanism, which involves hyperactivation of glutamate receptors, massive influx of  $Ca^{2+}$  cations resulting in "overactivation" of selected

signaling pathways. Different levels of magnitude and heterogeneity of neurodegeneration were observed following glutamate application in cultured neurons, however. Depending on the intensity of glutamatergic insult, apoptotic or necrotic types of cell death can occur. Intense glutamatergic stimulation causes necrosis, mediated by acute mitochondrial dysfunction, whereas milder glutamate influence results in the activation of several intracellular pathways, associated with other cell death types (Lau and Tymianski 2010).

The crucial player in glutamate-mediated excitotoxicity is a calcium influx into the cells, mainly through NMDA receptor channels (Lau and Tymianski 2010). NMDA receptors (dependent on subunit-specific differences) are approximately 3-4 times more permeable to  $Ca^{2+}$  than unedited  $Ca^{2+}$ -permeable AMPA, and Ca<sup>2+</sup>-permeable AMPA receptors still show a slightly greater Ca<sup>2+</sup> permeability than KA receptors (Traynelis et al. 2010). After three-phase time-dependent rise of calcium that includes initial 5-10-min-lasting acute phase, subsequent 2-h latent phase with normal Ca<sup>2+</sup> concentration followed by calcium sequestration in mitochondria in third sustained phase, a metabolic acidosis and free radical generation develop that result in mitochondrial damage. Glutamate-mediated calciumdependent neurodegeneration has also been linked to the generation of nitric oxide (NO for a nitric oxide synthase (NOS) knockout mice demonstrated attenuated NMDA-mediated excitotoxicity). More detailed studies revealed a structural scaffold – a postsynaptic density protein of molecular weight 95 kDa (PSD-95) connecting NR2B subunit of NMDA receptor and NOS. It is of particular importance for NO is directly neurotoxic, nitrosylating GADPH (glyceraldehyde 3-phosphate dehydrogenase, an ubiquitous glycolysis enzyme), which together with ubiquitin ligase (Siah1) enhances p300/CBP-associated acetylation of nuclear proteins (Lau and Tymianski 2010), including histones, thereby potentially influencing gene expression. Produced NO, together with free superoxide radicals, also causes protein modifications, lipid peroxidation, and DNA damage, resulting in subsequent cell death. Caspases, classical apoptotic cysteine proteases, activated downstream of calcium influx, and mitochondrial dysfunction play a significant role in this process as well. Other cellular cysteine proteases, calpains, might also be involved in glutamatergic excitoxicity by the cleavage and release of apoptosisinducing factor (AIF) from the mitochondria, which after nuclear translocation causes cell death, preceded by chromatin condensation and DNA fragmentation. Moreover, AIF release is linked to the activation of PARP-1 (poly(ADP)-ribose polymerase), enzyme involved in DNA repair in an alternative excitotoxic mechanism (Lau and Tymianski 2010).

The normal nontoxic Glu concentration between 1 and 3  $\mu$ M in the brain is maintained by excitatory amino acid transporters, mainly EAAT1 and EAAT2, predominantly expressed in nonmalignant astrocytes. However, as demonstrated in vitro, Glu concentration exceeding 500  $\mu$ M is present in cultured glioma cells. Likewise Glu concentrations in the range of 100  $\mu$ M were reported at the tumor margin of the glioblastoma-bearing patients (de Groot and Sontheimer 2011).

In contrast to neurons and glial cells, glioma cells release Glu mainly by employing  $x_c^-$  cysteine–Glu exchanger system, which requires extracellular

cysteine to be incorporated to the cell and intracellular Glu to be released outside the cell. It has been suggested that Glu may be generated from glutamine via a glutaminase reaction and/or by glutamate dehydrogenase (GDH). These notions are supported by the observation that high levels of liver type glutaminase and GDH have been demonstrated in gliomas. Their expression might be promoted by the oncogene *c-myc*, which stimulates glutamine metabolism and is overexpressed in treatment-resistant glioma stem cells. Additionally, EAAT1/2 transporters are often dysfunctional in glioma cells, and their expression (especially EAAT2) seems to be inversely correlated with the degree of malignancy with almost no expression in GBM cells (de Groot and Sontheimer 2011). In consequence glioma cells release high amounts of glutamate while having impaired glutamate uptake system, which results in excessive amounts of Glu in peritumoral area.

On the other hand, iGluRs and mGluRs are ubiquitously present on neurons and glial cells (Bowie 2008). Glu released from gliomas may then exert peritumoral excitotoxicity directed to neurons and oligodendrocytes. Astrocytes are usually resistant, even to high glutamate concentration (millimolar range). Still, peritumoral astrocytes probably lose their glutamate uptake potential and/or ability to synthesize glutamine from Glu. Thereby, high-grade gliomas probably also kill astrocytes, which is supported by observations that the space surrounding expansive tumors is devoid of normal brain cells (de Groot and Sontheimer 2011).

Another NMDA receptor agonist quinolinic acid (QA) might be considered as an additional endogenous neurotoxin involved in cancer-mediated neurotoxicity. OA is produced in kynurenine pathway, metabolizing an essential amino acid tryptophan (TRP). To achieve the final product, nicotinamide adenosine dinucleotide (NAD), yet another potential neurotoxin, 3-hydroxyanthranilic acid (HAA), and two endogenous neuroactive intermediates - the NMDA antagonist, kynurenic acid (KYNA), and the neuroprotectant, picolinic acid (PIC) – are generated along the pathway. The kynurenine pathway has been shown to be present in neurons, astrocytes, and microglial cells, as well as in brain tumor cells (Chen and Guillemin 2009). The differential pattern of kynurenine pathway expression was also demonstrated in neuroblastoma cells when compared to neurons. Neuroprotective PIC was produced primarily by immature and mature neurons, but not by the tumor cells. In contrast, neuroblastoma cells were able to produce the excitotoxin QA (Guillemin et al. 2007). QA is a heterocyclic amino acid that selectively activates the neuronal NMDA receptors and has been demonstrated to have immunomodulatory and neurotoxic properties, causing neuronal cell death after direct intracerebral administration (Chen and Guillemin 2009). It has been shown that QA can directly increase the proliferation rate of brain tumor cells in vitro (Guillemin et al. 2007) while being directly toxic to the surrounding neurons and thus mirroring the effects mediated by high amounts of glutamate. Interestingly, this effect can be abrogated by endogenous neuroprotectants, synthesized in the same pathway. KYNA and PIC may effectively block QA-induced neurotoxicity without affecting its excitatory effect. Kynurenic acid acts on the glycine modulatory site at low concentrations while at higher ones on the glutamate site of the NMDA receptors. KYNA was also shown to antagonize AMPA receptors and the alpha 7 nicotinic acetylcholine receptors.

Accordingly, KYNA was reported to exert sedative and anticonvulsant effects; it was also proved to be protective against brain ischemia. PIC is less potent and appears to act via a different mechanism attenuating calcium-dependent glutamate release and/ or chelating endogenous zinc (Chen and Guillemin 2009). Interestingly, neurons seem to synthesize it, while neuroblastoma cells do not (Guillemin et al. 2007).

Furthermore, quinolinic acid and 3-hydroxyanthranilic acid can effectively suppress T cell proliferation (Chen and Guillemin 2009) thus potentially attenuating the immune defense system. This could be mediated by overexpressed indoleamine 2,3-dioxygenase (IDO) enzyme, a component of the kynurenic pathway, involved in the conversion of tryptophan to *N*-formylkynurenine. IDO is being expressed by a plethora of cells, including glial tumor cells, dendritic cells, and monocytes in the tumor's vicinity, which in turn has implicated this molecule as one of the key mediators of the tumor immune escape. Beyond the tryptophan depletion, accumulation of IDO metabolites, 3-hydroxyanthranilic acid, KYNA, and QA, within the tumor environment and in the blood seems to increase the suppression of antitumor immune responses being quite detrimental in particular for the T cell, inducing both their anergy and death, as well as to impair NK cellular cytotoxicity by particularly targeting the expression of key NK receptors involved in target recognition (Zamanakou et al. 2007).

A commonly voiced concern about the possible role of QA in neuronal damage is that its concentrations in the blood and cerebrospinal fluid (CSF) are usually lower than these required to induce neuronal damage. However, the QA levels may be many times greater in the extracellular space around activated glial cells (Stone et al. 2012), or brain tumors, and there may be subsets of neurons which are particularly sensitive to quinolinic acid toxicity like neurons within the dentate gyrus of the hippocampus.

Notably, an enhanced degradation of TRP is observed in patients with various types of cancer. On the other hand patients with decreased TRP serum/plasma levels and increased concentrations of tryptophan catabolites have been reported to be involved in the development of mood disorders, especially depression. IDO activation, described above, leads to the accumulation of neurotoxic metabolites, which are supposed to induce depressive-like behavior. Resulting imbalance between neuroprotective (KYNA) and neurotoxic (QA) metabolites has been proposed to be critical for the development of depressive symptoms in patients with CNS tumors. On the other hand, decreased tryptophan availability might also affect neurosignaling by impaired synthesis of neurotransmitters serotonin and NAD, resulting in the development of depression (Kurz et al. 2011).

Taken all together, aberrant activation of the kynurenine pathway may contribute to the persistence and development of brain tumors and might be, at least in part, responsible for neurotoxicity associated with them.

## 3 Important Scientific Research

Direct evidence for neurotoxic function of glutamate released by glioma cells was provided for the first time in 1999. It was demonstrated then that established glioma

cell lines and primary cultures grown from the patient tissues harvested during biopsy release large amounts of glutamate, resulting in elevations of Glu that exceed 100  $\mu$ M within hours sufficient to activate glutamate receptors on hippocampal neurons in coculture experiments and to induce delayed Ca<sup>2+</sup>-dependent cell death (Ye and Sontheimer 1999).

Crucial data, which confirmed that glutamate excitotoxicity contributes to tumor expansion in vivo, were obtained by implanting different clones of rat glioma C6 cells in striata of adult rats. Clones that actively released glutamate developed aggressive tumors that expanded faster than tumors composed of wt C6 cells or derived from clones that were deficient in glutamate. Tumor size correlated with glutamate secretion by these clones and was not affected by proliferation rate and mitotic activity; as expected it also correlated with survival time of tumor-bearing animals. Consistent with the coculture experiments, tumors from clones releasing high amounts of Glu generated high degree of degenerated neurons, accompanied by the inflammatory response in surrounding brain areas (Takano et al. 2001). Based on this data it was concluded that glutamate acts as an important autocrine/ paracrine signal that promotes glioma cell invasion and the formation of tumor metastasis (Lyons et al. 2007).

Subsequent studies proved that GluR antagonists negatively affect proliferation of different types of cancer cells in vitro (Rzeski et al. 2001) and growth of implanted tumors in vivo (Stepulak et al. 2005; Takano et al. 2001). Accordingly, the involvement of glutamate receptors in the proliferation and migration of cancer cells was also suggested. Follow-up study by Ishiuchi et al. provided evidence that gliomas express functional AMPA receptors (Ishiuchi et al. 2002). NMDA and kainate receptor subunits were also demonstrated in brain tumor-derived cell lines (Stepulak et al. 2009). Still, their functional significance was not unequivocally proved. AMPA receptors seem to be crucial for promoting tumor cell proliferation and invasion. The presence of GluR2 subunit of the AMPAR is critical to the receptor function. In its edited form GluR2 makes AMPAR impermeable for calcium entry into the cell, resulting in attenuated ability of glioma cells to invade in vitro and in vivo, as demonstrated by overexpression of GluR2 subunit (Ishiuchi et al. 2002).  $Ca^{2+}$  imaging experiments showed also that the released glutamate activates AMPAR and induced intracellular Ca<sup>2+</sup> oscillations that are essential for cell migration (Lyons et al. 2007). Interestingly, brain-derived tumor cells expressed also unedited form of GluR2 subunit (Stepulak et al. 2009), thus enabling Ca<sup>2+</sup> influx through AMPAR functioning ion channel. A similar situation might take place in brain tumors in vivo; direct proof requires further studies, though.

The significance of AMPAR expression in brain tumors was also demonstrated in the clinic. Lack of GluR2 expression has been shown to be an independent prognostic factor for a poor prognosis in patients with high-grade glioma (Colman et al. 2010). GluR2 was absent in fast-growing GBM-derived tumor stem cells and high-grade glioma specimens but was expressed in slow-growing stem cells and low-grade glioma specimens (Beretta et al. 2009). Increased GluR1 expression was observed in glioblastoma samples when compared to anaplastic astrocytomas and low-grade gliomas. AMPA-mediated signaling in glioma cell lines was attributed to MAP kinase (ERK1/2) activation, and gene silencing of the GluR1 AMPA receptor subunit results in abrogation of AMPA-mediated signaling and tumor growth (de Groot et al. 2008). It has been also shown that GluR1 overexpression in glioma cells with subsequent activation of focal adhesion kinase (FAK) is responsible for increased cell adhesion to the extracellular matrix (ECM) in vitro (de Groot and Sontheimer 2011).

GluR2 overexpression in glioma cells inhibits proliferation and induces apoptosis by inactivating extracellular signal-regulated kinase (ERK)1/2 via Src kinase (Beretta et al. 2009). MAP kinase seems to be an intracellular mediator connecting mGluR3 receptor with bone morphogenetic proteins (BMP)-Smads transcription factors signaling. Activation of mGluR3 results in sustained undifferentiated state of glioma-initiating cells, whereas pharmacological blockade of mGluR3 stimulates differentiation of these cells and limits the growth of implanted tumors (Ciceroni et al. 2008). Akt kinase pathway, crucial for proliferation of various cancer cells types, seems to be another important downstream effector of AMPAR-mediated signaling in glioma cells. A dominant-negative form of Akt inhibits cell proliferation and migration accelerated by overexpression of  $Ca^{2+}$ permeable AMPA receptor. In contrast, introduction of a constitutively active form of Akt rescues tumor cells from apoptosis induced by the conversion of Ca<sup>2+</sup>permeable AMPA receptor to Ca<sup>2+</sup>-impermeable receptors by the delivery of GluR2 cDNA (Ishiuchi et al. 2007) proving the importance of this type of signaling for tumor cell homeostasis. Since glutamate receptors appear to be vital for glioma invasion and metastasis, inhibition of these receptors and/or GluR-dependent signaling pathways opens a new field for therapeutic intervention that should result in an inhibition of tumor invasion, along with reduction of neurotoxic side effects mediated by the tumor.

Expansion of the tumor may explain a frequent occurrence of seizures initiated by the tumor and the surrounding brain. Over 80 % of glioma patients suffer seizures during the course of their diseases (de Groot and Sontheimer 2011). However, the mechanism responsible for these seizures has not been unequivocally proved and may be related to the expression and activation of GluRs in different types of brain tumors and glial cells. Subunits from each receptor class, including ionotropic NR1, NR2A/B (NMDA) GluR1, GluR2 (AMPA), and GluR5-7 (kainate) and metabotropic mGluR1, mGluR2/3, mGluR5, and mGluR7a subtypes, were demonstrated in glioneuronal tumors from patients with intractable epilepsy. Group I mGluRs (mGluR1 and mGluR5) were highly represented, whereas low expression of mGluR2/3 (group II) and mGluR7 (group III) was observed in the neuronal component of the tumors, supporting the central role of glutamatergic transmission in the mechanisms underlying the intrinsic and high epileptogenicity of these lesions. Increased expression of mGluR2/3, mGluR5, and GluR5-7 proteins in peritumoral reactive astrocytes suggests its possible contribution to the generation and propagation of seizure activity (Aronica et al. 2001). In astrocytes iGluRs, especially AMPARs, are able to form functional ion channels, and their activation can alter astrocytes morphology and function, whereas mGluRs appear to regulate the expression of Glu transporters (de Groot and Sontheimer 2011). Astrocytes predominantly express mGluR3 and mGluR5, and their activation enhances glutamate uptake, thus protecting neurons from glutamate-mediated toxicity. These mechanisms are often disrupted under pathologic conditions. The mGluR5-mediated upregulation of glutamate transport is lost in chemically activated astrocytes in rat ALS model, and an enhanced release of glutamate induced by the activation of glial mGluR5 receptors contributing to glutamatemediated neuronal death was observed after status epilepticus (D'Antoni et al. 2008). Thus, tumor-released glutamate may potentiate excitotoxicity mediated by astrocytes by activating mGluR5 receptors. In microglia, activation of mGluR2 induces mitochondrial depolarization and apoptosis or, alternatively, induces an activated phenotype, which underlies microglia-mediated neurotoxicity, mediated probably by the secretion of TNFa and Fas ligand from activated microglia (D'Antoni et al. 2008).

Recently, the involvement of KARs activated by the endogenous agonist glutamate in the generation and propagation of epileptiform seizures was discussed (Vincent and Mulle 2009). Nonsynaptic source of glutamate derived from the activity of astrocytes increases the activation of KARs in neighboring interneurons. On the other hand, apoptotic and necrotic death of neurons is associated with dosedependent kainate (KA)-induced excitotoxicity in vivo (Vincent and Mulle 2009; Wu et al. 2005) and involvement of KA receptors in epileptiform seizure induction was demonstrated by injection of KA in rodents (Vincent and Mulle 2009). Thereby, it is possible that activation of KAR by glioma-released glutamate could contribute to the development of the seizures observed in glioblastoma patients. This hypothesis is supported by the observation that NMDA receptor antagonists given before or after the induction of KA-mediated seizures prevent most of the excitotoxic damage, suggesting pivotal role of NMDA receptors in KA-induced excitotoxicity (Vincent and Mulle 2009). Seizures might be also activated by quinolinic acid (QA) acting on NMDA receptors and producing neuronal damage (Stone et al. 2012), which have been shown to be produced by neuroblastoma cells (Guillemin et al. 2007).

## 4 Recent Findings

Despite recent improvements in treatment strategies, the results of therapy in patients with glioblastoma are unsatisfactory. Effective treatment is limited due to biological properties of the tumor and the clinical advancement of the disease upon diagnosis. Because of the low efficacy of the traditional chemotherapy, new therapeutic tools acting selectively on specific metabolic pathways pathologically deregulated in glioma cells are still needed. On the other hand, development of new tools attenuating tumor- and glutamate-induced neurotoxicity is of great importance.

Since, new types of the targeted therapy, including blockade of growth factors, like EGFR, GFs-mediated downstream signaling (Ras–ERK1/2, Akt–mTOR, PKC), proteasome, and microtubule inhibitors, proved to be of limited value in the clinic (Adamson et al. 2009), glutamatergic signaling system has attracted great

attention. As described above, glutamate receptor signaling has been implicated in tumor biology, and receptor antagonists have been shown to be sufficient to suppress cancer growth in experimental studies (Arcella et al. 2005). Unfortunately, the majority of tested compounds, although promising in animal experiments, failed in clinical trials (Lau and Tymianski 2010). It is of interest, however, that in contrast to other NMDA receptor antagonists, memantine has been approved for the treatment of Alzheimer disease and has shown some antitumor effects in vitro (Rzeski et al. 2001; Takano et al. 2001). It is worth mentioning that doses, which inhibit tumor cells proliferation, are much higher than those applicable for Alzheimer disease. The other iGluR antagonists show similar properties, even though it has been shown that some of them exert their activities at relatively low, clinically relevant doses both in vitro (Rzeski et al. 2001) and in vivo (Stepulak et al. 2005). Unfortunately, potential side effects, psychotic in particular, are limiting factors of iGluR antagonists' therapy (Lau and Tymianski 2010).

On the other hand, very promising results of treatment of newly diagnosed glioblastoma patients with talampanel combined with temozolomide (TMZ) and radiation therapy (RT) were reported. Talampanel, an oral, noncompetitive AMPAR antagonist with excellent brain penetration, combined with RT and TMZ, significantly prolonged median survival time, without any additional toxicity, as compared to RT–TMZ-treated patients (Grossman et al. 2009). This provides a solid base for future studies with similar, newly synthesized inhibitors.

Recently, another group of potential therapeutic agents that inhibit glutamate release has draw attention. Glioma cells in the presence of deficient glutamate uptake systems upregulate the glutamate exchanger  $X_c$ - cysteine–glutamate antiporter that leads to progressive Glu accumulation in peritumoral tissue (de Groot and Sontheimer 2011). A disruption of this system using shRNA attenuated Glu release in vivo, prolonged survival of tumor-bearing animals, and diminished glioma-induced neurodegeneration and brain edema (Savaskan et al. 2008). It has been shown that an inhibition of the  $X_c$ - transporter by sulfasalazine results in tumor volume reduction by 60–80 % in animal model (Sontheimer 2008). Despite several limitations such us poor systemic bioavailability because of its rapid cleavage by colonic bacteria and a half-life of ~80 min in circulation, clinical trials with sulfasalazine in the treatment of newly diagnosed low-grade gliomas are ongoing (de Groot and Sontheimer 2011).

Another drug aimed on inhibition of glutamate release and attenuation of its neurotoxicity is riluzole (2-amino-6-trifluoromethoxybenzothiazole). Riluzole is an anticonvulsant and neuroprotective agent approved by FDA for different indications (treatment of amyotrophic lateral sclerosis – ALS). Its main mechanisms of action involve inhibition of Na<sup>+</sup>-gated channels, postsynaptic blockade of ionotropic GluRs without direct receptor interaction, and potentiation of glutamate uptake by activation of glutamate transporters (Bellingham 2011).

Riluzole has been shown to inhibit proliferation and glutamate release in melanoma, both in vitro and in vivo (Namkoong et al. 2007); however, it has not been tested in glioma treatment, yet. Nevertheless, inhibition of cellular systems of glutamate release seems to be an interesting target for future therapies of brain tumors aimed on reduction of tumor-mediated neurotoxicity.

On top of tumor-mediated neurotoxicity, one should remember that present therapeutic strategies result in significant toxicity directed to brain cells (Rinne et al. 2012). Thus, the development of new neuroprotective agents has become a necessity. One of the most promising approaches involves application of a novel class of drugs aimed at disruption of protein complexes between GluRs and scaffolding proteins. It is of particular value that these specific peptides disrupt interactions between NMDAR and PSD95 or KAR and PSD95 without influencing synaptic activity or calcium influx, which in turn inhibits downstream neurotoxic signaling without interfering with neurotransmission (Vincent and Mulle 2009). Another group of peptides, named membrane-associated guanylate kinase (MAGUK) inhibitors, dissociates spatial relationship between NMDAR and NOS in a competitive manner, inhibiting NOS-mediated neurotoxicity. One of these peptides – NA1 – is currently in a phase II clinical trial (Lau and Tymianski 2010).

For years, agents such as psychostimulants and acetylcholinesterase inhibitors have been used to treat cognitive dysfunction in patients with brain tumors. Recently, however an improvement in cognitive functions in high-grade glioma patients treated with methylphenidate and modafinil, known for their neuroprotective capabilities (van Vliet et al. 2008; Ludolph et al. 2006), was reported. Donepezil, another compound with proven neuroprotective activity against glutamate-induced excitotoxicity (Shen et al. 2010), has also been shown to improve cognitive performance in patients with brain tumors.

Currently clinical trials are being conducted in order to determine the potential neuroprotective effects of memantine and lithium in patients with brain metastases (Correa 2010). Finally, there is some evidence that GluR5-containing KARs represent a novel target for AED (antiepileptic drug) development, since GluR5 antagonists prevent and interrupt limbic seizures without overt side effects in the experimental models (Vincent and Mulle 2009), thereby are of potential interest also for glioma-mediated seizures.

#### 5 Conclusion

Brain tumors, especially glioblastoma, are still a therapeutic conundrum in the context of short survival times and low quality of life of the patients that arise from the neurotoxicity mediated by both tumor and side effects of applied therapies. An improvement of anticancer therapy on one side and attenuation of neurotoxicity on the other define future challenges on the field. Thereby, additional therapeutic approaches are needed to supplement existing strategies (Candolfi et al. 2009). A deregulated glutamatergic system in glioma is definitely one of the plausible therapeutic targets. Still, a number of questions regarding the role of ionotropic glutamate receptors in carcinogenesis remain unanswered. Even though initial reports confirmed the role of glutamate and its receptors in oncogenesis and their antagonists proved to exhibit a therapeutic potential, a practical implementation of those compounds remains elusive. The main obstacle relates to their serious side effects, in particular related to the CNS. Moreover, relevant information on the role

of the aforementioned receptors in peripheral tissues is also missing. The receptors themselves exist in various configurations of subunits that affect their pharmacological proprieties; on top of that receptor subunits undergo posttranslational and epigenetic modifications, which additionally increase their variability. iGluR antagonists are usually receptor or even subunit specific; accordingly a prospective therapy will probably require prior analysis of receptors in a given tumor. None-theless, existing data, in particular the results of clinical trials with talampanel in gliomas, are encouraging and support the search for novel antagonists of ionotropic glutamate receptors with superior pharmacologic proprieties to be implemented into antitumor and neuroprotective therapeutic protocols. On the other hand, strategies aimed on the inhibition of glutamate release with subsequent disruption of autocrine glutamate signaling loop, or strategies involving intracellular targets, where different signaling pathways converge, including those implementing histone deacetylase inhibitors (HDI) together with other standard or targeted types of therapy, are also of interest.

Another important advance that has recently changed our way of thinking about glial tumors biology relates to the concept of the so-called tumor stem or tumorinitiating cells (TSCs). It has been recently proven that within a total population of glioblastoma cells, there appears to be a small subpopulation of cells that are highly tumorigenic, with capacity for self-renewal (Singh et al. 2004). Because they share many properties with neural stem cells, it has been suggested that they originate from neuronal stem cells (Alcantara Llaguno et al. 2009). These cells are functionally defined with self-renewal and cell differentiation in vitro and their capability for tumor propagation in vivo (Li et al. 2009). The most important aspect of their presence within glial tumors is that this population appears to be particularly resistant to conventional radiation (Bao et al. 2006) and chemotherapy (Liu et al. 2006). What is of particular interest, however, is the fact that only recently this specific cellular subpopulation has been proven to express high concentrations of functional calciumpermeable AMPA receptors when compared to the differentiated tumor cultures consisting of non-stem cells. This raise the possibility that glutamate secretion in the glioblastoma microenvironment may stimulate brain tumor-derived cancer stem cells (Oh et al. 2012). This finding suggests that therapeutic strategies aimed at glutamatedependent excitotoxic pathway disruption might also influence TSCs, possibly initiating their differentiation thus rendering them more susceptible to conventional therapeutic modalities. This hypothesis requires further substantiation, though.

A possible strategy to overcome these obstacles involves more personalized diagnosis and treatment, including analysis of genetic abnormalities of particular tumor cell populations, as well as assessment of potential treatment efficacies of arrays of drugs tested on primary cell cultures, derived from samples of tumors harvested during surgery.

In the context of neurotoxicity approaches that render physiological glutamatergic transmission intact and simultaneously disrupt excitotoxic pathways are of particular interest. Another potential therapeutic strategy for neuroprotection involves mimicry of the glutamate blocking activity of endogenous neuroprotective agents, such as kynurenic acid or melatonin. Nevertheless, a lack of significant success with therapeutic protocols that interfere with NMDA–AMPA-dependent mechanisms of excitotoxicity implies that some other potential therapeutic targets have to be explored.

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# Drug Treatments for Alzheimer's Disease: Hopes and Challenges

Nady Braidy, Anne Poljak, Tharusha Jayasena, and Perminder Sachdev

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#### P. Sachdev

N. Braidy (🖂) • T. Jayasena

Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

e-mail: n.braidy@unsw.edu.au; t.jayasena@student.unsw.edu.au

A. Poljak

School of Medical Sciences, Faculty of Medicine and Bioanalytical Mass Spectrometry Facility, University of New South Wales, Sydney, Australia e-mail: a.poljak@unsw.edu.au

Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

Neuropsychiatric Institute, Prince of Wales Hospital, Sydney, Australia e-mail: P.Sachdev@unsw.edu.au

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#### Abstract

At present, no effective cure is available to slow down or prevent progressive neuronal loss in the pathogenesis of Alzheimer's disease (AD). All currently approved therapeutic agents provide symptomatic relief only. However, current agents under development exhibit potential protective effects on disease course. In this chapter, we consider currently marketed drugs for AD, including acetylcholinesterase inhibitors (AChEI) and antagonists of *N*-methyl-D-aspartate (NMDA) receptors. This work also discusses potential cognitive enhancers under clinical trial. We also review new therapies that are likely to slow down disease progression by mechanisms unrelated to amyloid or tau.

### Keywords

ACE inhibitors • Alzheimer's disease • Amyloid • Antioxidants • Tau

# 1 Introduction

Alzheimer's disease (AD) is an irreversible, progressive neurodegenerative disorder characterized by impaired learning and memory loss. One of the major pathological hallmarks of AD is the presence of a high density of extracellular senile plaques containing amyloid beta (A $\beta$ ) in the hippocampus and cerebral cortex (Shen et al. 2008). Irregular processing and accumulation of A $\beta$  can impact negatively on important metabolic pathways within neurons and glial cells leading to neuronal and synaptic dysfunction and neurodegeneration. A $\beta$  generation is regulated by the sequential proteolytic activity of  $\beta$ -secretase (BACE) and  $\gamma$ -secretase proteases (De Strooper et al. 2010; Hashimoto et al. 2010; Marks and Berg 2010; Neugroschl and Sano 2010; Tabaton et al. 2010). When the  $\alpha$ -secretase pathway is used for the processing of amyloid precursor protein, no A $\beta$  is generated (Nord et al. 2010). Presenilins 1 and 2 are the core components for the catalytic activity of  $\gamma$ -secretases, which are necessary for A $\beta$  formation and progression of AD (Rocher-Ros et al. 2010).

Intracellular neurofibrillary tangles (NFTs) containing hyperphosphorylated tau protein are the second hallmark for AD (Attems et al. 2012; Wei et al. 2012). In the healthy neuronal population, tau protein helps to stabilize microtubules in the cytoskeleton (Kon et al. 2012). Although the mechanism for tau hyperphosphorylation in AD remains unclear, this hyperphosphorylation also occurs in numerous dementias and in the pathogenesis of other neurodegenerative disorders including Parkinson's disease (PD) (Lasagna-Reeves et al. 2012).

Prodromal mild cognitive impairment precedes AD, with more than half of these individuals progressing to AD within 4 years of symptomatic onset. Most individuals with mild cognitive impairment show early signs of AD pathology in their brain, including enhanced A $\beta$  deposition and NFTs (Petersen et al. 1999). Owing to the postmitotic nature of neuronal cells, therapeutic agents that can slow down and/or reverse the pathogenic AD cascade prior to irreversible neuronal damage may alter the disease course in AD (Olazaran et al. 2010).

AD is the most common cause of dementia among elderly people over 65 years of age. For every 5-year age group beyond 65, the percentage of people with AD doubles. Scientists estimate that around 4.5 million people worldwide suffer from AD (Olazaran et al. 2010). Despite the alarming increase in the prevalence of AD, there are no disease-modifying drugs available. Currently approved treatments include the acetylcholinesterase inhibitors (AChEI) donepezil, galantamine, and rivastigmine and memantine, an antagonist of *N*-methyl-D-aspartate (NMDA) receptors. Other proposed therapies include  $\beta$ - and  $\gamma$ -secretase inhibitors, glucagon-like peptide-1 (GLP-1), bapineuzumab, solanezumab, statins, nonsteroidal anti-inflammatory drugs (NSAIDs), Cerebrolysin, psychotropic agents, and dietary supplements, vitamins, and nutraceuticals.

# 2 Acetylcholinesterase Inhibitors

The progressive degeneration of cholinergic neurons in the basal forebrain and reduced cholinergic neurotransmission in the cerebral cortex and other areas contribute significantly to impaired learning and memory observed in patients with AD. This neurochemical abnormality has provided the rationale for the development of first-line drugs for the treatment of AD which induce the cholinergic system in which the neurotransmitter acetylcholine is lacking (Pepeu and Giovannini 2009). Acetylcholinesterase inhibitors (AChEIs) are the first class of agents approved by the US Food and Drug Administration (FDA) for the treatment of AD. Tacrine (Cognex) was first approved in 1993 but is now rarely used due to elevations in hepatic enzymes and will not be discussed here. In 1996, donepezil was approved under the trade names Aricept, Adonep, Yasnal, and others. It was followed by rivastigmine (Exelon, Prometax, Ristidic, Vastigmex, or Evertas) in 2000 and galantamine (Reminyl, Razadyne, or Nivalin) (Pohanka 2011).

Inhibitors of acetylcholinesterase (AChE) share similar mechanisms of action while exhibiting additional pharmacological properties. The response to treatment with AChEIs is similar, and no individual AChEI has been shown to display superiority to others in efficacy studies. AChEIs bind to the synaptic cleft, thus stimulating the release of acetylcholine from the presynaptic cholinergic terminal and increasing the potential for interaction with the postsynaptic cholinergic receptor, therefore promoting the function of the cholinergic receptor (Pohanka 2011). Donepezil is a selective reversible AChEI with an affinity to AChE which is 1,000-fold higher than towards butyrylcholinesterase (BCHE) (Perdomo et al. 2003).

	AChEIs		
Characteristics	Donepezil	Rivastigmine	Galantamine
Indications	Mild to moderate and severe AD	Mild to moderate AD; Parkinson's disease dementia	Mild to moderate AD
Half-life (hrs)	70	1.5	7
Administration regime	q.d.	b.i.d. for capsule; q.d. for the patch	b.i.d. for the non-ER form; q.d. for the ER form
Metabolism by hepatic CYP enzymes	2D6, 3A4	No	2D6, 3A4
Protein binding (%)	96	40	18
Bioavailability	100	40	90
Absorption delayed by food (hrs)	No	No	Yes (1–2.5)
Time to reach peak serum concentration (hrs)	3–4	1	1 (2.5 with food); 4.5 with ER form
Titration schedule	Initial dose of 5 mg then advance to 10 mg after 1 month	Oral form: 1.5 mg b.i.d. for 4 weeks then 3 mg for 4 weeks then to 6 mg b.i.d. if tolerated	Non-ER form: 4 mg b.i.d. then 8 mg b.i.d. after 1 month to 12 mg b.i.d. after another month
		Patch form: initial patch of $5 \text{ cm}^2$ for 1 month then the $10 \text{ cm}^2$ patch	ER form: 8 mg q.d. after 1 month then to 16 mg for 1 month then 24 mg q.d. after another month
Adverse effects	Gastrointestinal, nausea, vomiting	Gastrointestinal, nausea, vomiting	Gastrointestinal, nausea, vomiting
	Reduced heart rate, sleep disturbances, muscle cramps	Reduced heart rate	Reduced heart rate

Table 1 Pharmacodynamics of AChEIs

Abbreviations: q.d. every day; b.i.d. twice a day; ER extended release

Rivastigmine is a nonselective mixed inhibitor of both AChE and BCHE (Naik et al. 2009). Galantamine is an allosteric modulator of AChE with limited affinity towards BCHE. Apart from interaction with AChE, galantamine neuroprotective effects can be further attributed to its allosteric modulation of nicotinic acetylcholine receptor (nAChR) (Feng et al. 2010). The basic pharmacology and pharmacodynamics of these drugs are summarized in Table 1.

It remains unclear as to whether AChEIs have any disease-modifying properties in AD. It has been suggested that enhanced cholinergic transmission can slow down A $\beta$  production (Kimura et al. 2005). Several imaging studies have shown a marked reduction in brain atrophy in patients treated with AChEIs (Krishnan et al. 2003). One study

has shown reduced decline in cognitive function in patients treated with AChEIs or in combination with memantine compared to placebo (Lopez et al. 2009). However, no disease-modifying effects have been reported in another study (Jack et al. 2008).

Huperzine, a metabolite in the Chinese lycopod *Huperzia serrata* (Huperziaceae), is another drug that has been shown to have a neuroprotective effect suitable for the treatment of AD. It exists in two forms: A and B. However, the A variant is most likely pharmacologically relevant. It is a noncompetitive inhibitor of AChE, with no effect on BCHE. As well, it is a potent antagonist of the NMDA receptor (Gao et al. 2009). It can also attenuate oxidative stress and upregulate endogenous antioxidant activity in the CNS (Coleman et al. 2008). Huperzine has become target for extensive pharmacological research and is available for the treatment of neurodegenerative diseases in China. However, it is yet to be approved in the USA, Europe, and Australia. Favorable outcomes from a current clinical trial will allow the drug to be approved for therapy purposes in those countries (Ma et al. 2007).

#### 3 Memantine

Memantine antagonizes the NMDA receptor by replacing the potassium in the NMDA ion channel to reduce the efflux of intracellular calcium into neurons and prevent excitotoxicity and subsequent apoptosis. Prior to its approval for the treatment of AD, memantine was used for the treatment of Parkinson's disease in Germany (Winblad et al. 1999). In 2004, memantine (Namenda, Ebixa, or Axura) was approved for the treatment of moderate to severe AD by the FDA in the USA (Ferris et al. 2003; Moebius et al. 2003a, b; Reisberg et al. 2003) (Table 2).

Whether the neuroprotective effects of memantine on neuronal cells observed in vitro are responsible for the symptomatic relief observed in clinical trials remains to be examined. Reduced A $\beta$  deposition and tau hyperphosphorylation have been reported in in vitro studies after treatment with memantine (Martinez-Coria et al. 2010). Memantine has also been shown to provide beneficial effects on long-term potentiation (LTP), which correlates with memory function (Frankiewicz and Parsons 1999). This observation provides a plausible explanation for the symptomatic relief reported in patients undergoing therapy with memantine.

Memantine may be administered alone or in combination with an AChEI. No adverse effects have been reported with patients receiving both drugs. Pharmacotherapy with an AChEI and memantine continues until the patient either succumbs to AD or the quality of life of the patient is significantly compromised that the combination therapy is vain. Some long-term open-labeled studies have shown that combination therapy with an AChEI and memantine may attenuate the progression of AD, although other studies show conflicting results (Lopez et al. 2009; Schneider et al. 2009).

Indications	Moderate to severe AD
Half-life (hrs)	60–80
Metabolism by hepatic CYP enzymes	No
Protein binding (%)	50
Absorption delayed by food	No
Time to reach peak serum concentration (hrs)	3–7
Titration schedule	Initially 5 mg q.d. for 1 week, 5 mg b.i.d. for the next week, 10 mg in the morning and 5 mg in the evening for 1 week, and 10 mg hereafter

Table 2 Pharmacodynamics of memantine

Abbreviations: q.d. every day; b.i.d. twice a day

# **4** β-Secretase Inhibitors

 $\beta$ -Secretase cleaves APP into two peptide fragments, A $\beta$ 40 and A $\beta$ 42 (Zhou et al. 2012). Therefore, inhibition of  $\beta$ -secretase may reduce plaque formation and slow down the progression of AD. Preliminary studies using  $\beta$ -secretase knockout models showed only minor phenotypic changes, suggesting that  $\beta$ -secretase inhibitors may be an effective therapeutic strategy (Tang and Ghosh 2011). Novel  $\beta$ -secretase inhibitors have been developed as potential preventative and/or intervention therapy for AD.

Treatment with the  $\beta$ -secretase inhibitor (GRL-8234) has been shown to attenuate age-mediated decline in cognitive function in a transgenic mice model expressing APP after long-term inhibitor treatment (Chang et al. 2011). Moreover, the same study showed that GRL-8234 reduced brain A $\beta$  plaque load in these mice, suggesting a potential treatment for patients with late-stage AD (Chang et al. 2011).

TAK-070, a non-peptidic compound, has also been demonstrated to be a potent  $\beta$ -secretase inhibitor (Fukumoto et al. 2010; Takahashi et al. 2010). This compound can reduce the levels of soluble A $\beta$ , promote neurotrophic levels of APP, and inhibit the extracellular deposition of insoluble A $\beta$ . A clinical trial using TAK-070 is currently under consideration. Additional efficacy studies in rodents are underway (Fukumoto et al. 2010; Takahashi et al. 2010).

# **5** γ-Secretase Inhibitors

Inhibition of  $\gamma$ -secretase may also represent an important process to prevent the proteolytic processing of APP and regulate A $\beta$  formation (De Strooper and Annaert 2010). However, these inhibitors have been shown to cause adverse effects in preclinical animal models and human clinical trials. In one study, inhibition of  $\gamma$ -secretase induced embryonic toxicity, likely due to impaired activation of the notch, which is vital for cellular development and differentiation (Donoviel et al. 1999).

In a Phase II clinical trial, the  $\gamma$ -secretase inhibitor LY450139 leads to unwanted side effects including drug-related rashes and lightening of hair color with none

reported in the placebo group (Fleisher et al. 2008). Similarly, a Phase III trial for the  $\gamma$ -secretase inhibitor semagacestat was halted after the cognitive symptoms worsened in patients receiving the drug compared those who were given the placebo. Moreover, the incidence of skin cancers was increased in patients receiving semagacestat (Samson 2010). Therefore, the key question is whether undesirable side effects may result from the inhibition of cleavage of additional substrates for APP processing and whether a therapeutic window is necessary for the administration of these inhibitors to reduce A $\beta$  deposition.

Several agents, including NSAIDs and other organic compounds, have been recently shown to regulate  $\gamma$ -secretase activity, without interfering with notch signalling (Imbimbo and Giardina 2011; Imbimbo et al. 2011). These  $\gamma$ -secretase modulators may be efficacious for the treatment of AD without the serious adverse effects.

# 6 Bapineuzumab

Active immunization with synthetic full-length A $\beta$  peptide (AN1792) has been discontinued due to significant cognitive and neurological impairment occurring as a result of meningoencephalitis occurring in 6 % of patients (Orgogozo et al. 2003). The antibody-mediated response to AN1792 has been associated with the N-terminal residues (Lee et al. 2005). Subsequent studies using antibodies which target the N-terminus of A $\beta$  which do not contain T-cell activating epitopes can be protective against neurotoxicity and inhibit fibrillogenesis (McLaurin et al. 2002).

Bapineuzumab (AAB-001) is a humanized monoclonal antibody directed against the N-terminal region of A $\beta$ . Magnetic resonance imaging (MRI) abnormalities have been noted in a Phase I trial which generally resolve. There are also reports of vasogenic edema, likely due to an interaction of bapineuzumab with A $\beta$  occurring in the blood vasculature. These cases are largely associated with microhemorrhage which is asymptomatic (Black et al. 2010).

Additionally, bapineuzumab can improve cognitive performance assessed using the mini-mental state examination at lower doses compared to the placebo, while no change is observed in higher doses (>5 mg/kg). In a Phase II study, CSF t-tau and p-tau levels were significantly reduced at the end of the study compared with baseline in bapineuzumab-treated patients, while no change was detected in the placebo group. Interestingly, no difference was observed in A $\beta$  levels in either group (Blennow et al. 2012; Winblad et al. 2012). These studies highlight the downstream effects of bapineuzumab on the neurodegenerative process in AD, which may prove useful as a disease-modifying therapy in the early stages of AD.

# 7 Solanezumab

Solanezumab (LY2062430) is a humanized monoclonal antibody that targets the central region of A $\beta$  (Lachno et al. 2011; Reichert 2011; Samadi and Sultzer 2011).

Preclinical studies have shown that acute and subchronic treatment with solanezumab can ameliorate or reverse memory deficits in transgenic mice (Siemers et al. 2010; Panza et al. 2012; Uenaka et al. 2012). Unlike other disease-modifying immunotherapies, no effects on incidence or severity of cerebral amyloid angiopathy-associated microhemorrhages, a serious adverse effect associated with bapineuzumab, have been reported. Solanezumab displayed a good safety profile in a Phase II clinical study for AD and can significantly reduce cerebrospinal and plasma levels of  $A\beta$  and tau (Carlson et al. 2011; Farlow et al. 2012; Imbimbo et al. 2012).

In a Phase III trial, primary endpoints, both cognitive and functional, were not met in patients with mild to moderate AD. However, a statistically significant slowing of cognitive decline was shown in a prespecified secondary analysis of pooled data in patients with mild AD. Although solanezumab may be the first passive vaccine for the prevention of AD, immense scepticism still exists regarding the ability of this drug to slow down the progressive degeneration in patients with AD (Prodouz 2012). The next steps for solanezumab will be determined after discussions with pharmaceutical regulators.

### 8 Statins

Conflicting reports are available regarding the role of statins in improving brain function in AD. Epidemiological studies have shown a reduction in the risk of developing AD in patients using statins compared to those that have not been treated with cholesterol-lowering drugs. However, there are also numerous reports indicating significant cognitive decline following treatment with statins (Haag et al. 2009). Therefore, the usefulness of these drugs for the treatment of AD remains uncertain.

The neuroprotective effects of statins may be due to several mechanisms. Statins have demonstrated beneficial effects in the AD brain by enhancing degradation of extracellular A $\beta$  by inducing the release of insulin-degrading enzymes from the resident microglial cells. Increase levels of soluble A $\beta$  have been reported in microglia-depleted mice (Tamboli et al. 2010). However, since increased cholesterol levels is a risk factor for AD, it is difficult to determine whether the beneficial effects of statins in AD are related to the action of the statin or the improved lipid profile.

Numerous studies have been performed to determine the role of lipid-lowering agents for the treatment of AD. The Lipitor's Effect in Alzheimer's Dementia study found no significant difference in cognition or global function in patients with mild to moderate AD receiving Atorvastatin 80 mg/day for 72 weeks (Feldman et al. 2010). The Cholesterol Lowering Agent to Slow Progression of AD is a Phase III study currently underway to examine the safety and effectiveness of simvastatin to slow the progression of AD (McGuinness et al. 2010). Additional studies are needed to clarify the risks and benefits of statin therapy for the treatment of AD.

### 9 NSAIDs

A retrospective human epidemiological study showed that long-term use of NSAIDs, particularly ibuprofen, can protect against AD (Vlad et al. 2008). Similarly, experiments using rodent models for AD have predicted the effects of NSAIDs in humans, although little is known regarding the effect of NSAIDs on AD risk and pathogenesis. Prophylactic treatment of young 3xTg-AD mice with ibuprofen reduced intraneuronal A $\beta$  oligomers and hyperphosphorylated tau and enhanced cognitive function (McKee et al. 2008). Ibuprofen has also been shown to reduce reactive oxygen species production by inhibiting the activity of nicotinamide adenine dinucleotide phosphate oxidase (Wilkinson et al. 2012). NSAIDs represent a promising therapeutic target also due to their modulatory effect on  $\gamma$ -secretase activity, without inhibiting important  $\gamma$ -secretase-dependent signalling pathways (Pettersson et al. 2011).

#### 10 Glucagon-Like Peptide

Type 2 diabetes is a major risk factor for AD. Impaired insulin signalling appears to be a converging biochemical phenomena present in both diseases (Gotz et al. 2009; Ke et al. 2009). Apart from the pathological hallmarks in the AD brain, insulin receptors are reported to be desensitized in the AD brain, and this is referred to as type 3 diabetes (Steen et al. 2005). Therefore, effective treatment regime against type 2 diabetes may be of value in AD.

Activation of the glucagon-like peptide-1 (GLP-1) receptor has been recently shown to reduce A $\beta$  deposition and downstream toxicity in both cellular and rodent models for AD (Li et al. 2010). Moreover, GLP-1 has been shown to reduce neuronal APP levels in vitro (Perry et al. 2003). Taken together, these studies provide further support for the association between diabetes and AD.

### 11 Cerebrolysin

Cerebrolysin is a parenterally administered digested peptide preparation present in the pig brain and mimics the action of endogenous neurotrophic factors. This compound has demonstrated neuroprotective effects. Positive effects of Cerebrolysin on A $\beta$ - and tau-related pathologies, neuroinflammation, neurotrophic factors, oxidative stress, excitotoxicity, neurotransmission, brain metabolism, neuroplasticity, neuronal apoptosis and degeneration, neurogenesis, and cognition were demonstrated in experimental conditions (Allegri and Guekht 2012; Heiss et al. 2012; Masliah and Diez-Tejedor 2012; Menon et al. 2012; Sharma et al. 2012a, b; Thome and Doppler 2012; Vazquez-Roque et al. 2012; Xiao et al. 2012). As well, numerous studies have demonstrated symptomatic benefits in AD (Okamura et al. 2008). As a result, this drug is widely used in several countries, although more definitive trials are needed to ensure the effectiveness of Cerebrolysin as a monotherapy or in conjunction with other treatment regimens.

# 12 Psychotropic Agents

Behavioral disturbances are common in AD and include depression, agitation, irritability, abnormal motor behaviors, and psychosis (Cummings 2003; Porter et al. 2003). At present, no drug is currently available to treat these behavioral disturbances. The use of conventional and atypical antipsychotics can increase the risk of mortality from  $\sim 2.6$  % to  $\sim 4.5$  % in patients with AD (Kales et al. 2007).

Several 1–12-week-long clinical trials have shown that risperidone is efficacious in reducing psychosis in nursing home patients with AD at low doses (Ballard and Howard 2006; Ballard and Waite 2006; Fossey et al. 2006). Similarly, there is no evidence for the use of antidepressants in AD (Rosenberg et al. 2010; Weintraub et al. 2010). Several studies have shown that valproate has no benefit for the treatment of agitation compared to placebo and induces severe toxicity (Herrmann and Lanctot 2007; Herrmann et al. 2007). Anxiolytics should be avoided in patients with AD as they induce confusion. Short-term benzodiazepines such as lorazepam or clonazepam may be useful in patients with frequent episodes of agitation (Tariot et al. 1998). Given the current evidence for benefit and harm for the indication of antipsychotics in AD, these drugs should only be prescribed with caution to patients where non-pharmacological therapeutics have failed and their behavioral change is extreme, administering the treatment only when needed and informing the patient and responsible carer of the associated risks.

# 13 Dietary Supplements, Vitamins, and Nutraceuticals

Dietary supplements and vitamins have been used in clinical trials for AD. Nutraceuticals describe nontraditional compounds that display pharmaceutical effects but cannot be advertised for a health disease claim. Formulations of these substances vary, and no regulatory standards are available to verify the potency of these compounds except that these products contain the active substance advertised.

# 13.1 Docosahexaenoic Acid

Docosahexaenoic acid (DHA) is the main constituent of the cellular membrane in the CNS. The levels of DHA are reduced with age but can be replenished to normal levels following dietary supplementation. DHA has also been shown to reduce  $A\beta$  deposition in mice and improve cognitive performance when administered as a dietary supplement (Geleijnse et al. 2012; Morris 2012; Rosales-Corral et al. 2012; Stough et al. 2012). One trial reported improvements in episodic

memory after post hoc analyses in older individuals who have cognitive test scores below the normal threshold for young adults (Okamura et al. 2008). However, DHA supplementation did not alter cognitive function in another study (Quinn et al. 2010).

# 13.2 B Vitamins

Increased level of homocysteine in the blood is a major risk factor for cardiovascular disease, AD, and other dementia-related disorders and can potentiate the neurotoxic effect of A $\beta$  peptides in vitro. Homocysteine levels can be reduced by at least 30 % using regimens of B vitamins (B12 1 mg/day, B6 25 mg/day, and folic acid 5 mg/day). One trial showed a substantial reduction in homocysteine levels with these vitamins but did not reduce the rate of cognitive deterioration in AD patients compared to the placebo (Aisen et al. 2008).

# 13.3 Vitamin E

The antioxidant effects of vitamin E might be useful for the treatment of AD. In one study, vitamin E (1,000 IU b.i.d.) combined with selegiline (5 mg b.i.d.) delayed the time until patients with moderate AD require nursing home placement or succumb to the disorder, although no effect on cognition was reported (Sano et al. 1997). Another study reported no significant delay in conversion to AD or dementia when vitamin E was administered alone or with donepezil in patients with mild cognitive impairment (Petersen et al. 2005).

# 13.4 Homotaurine

Homotaurine (Vivimind) is an inhibitor of A $\beta$  aggregation that is naturally found in certain seaweeds. It is marketed in Canada as a nutraceutical with no specific health claims. Although a Phase II safety trial reported reduced CSF A $\beta$  levels, two follow-up 18-month Phase III trials did not show any efficacy (Aisen et al. 2006, 2011).

# 13.5 Ginkgo biloba

*Ginkgo biloba* leaves and extracts are currently used as over-the-counter preparations for the treatment of numerous medical conditions. The rationale for the use of *Ginkgo biloba* for the treatment of AD stems from its free radical scavenging activity or the inhibition of pro-inflammatory pathways or the potentiation of the cell stress response (Uluoglu and Guney 2010; Walesiuk and Braszko 2010; Serby et al. 2011). However, clinical studies are yet to validate the beneficial effects of *Ginkgo biloba* in improving cognitive function in AD patients.

#### 13.6 Resveratrol

Resveratrol is a naturally occurring phytochemical that is found in highest levels in red wine and the skin of red grapes. A recent epidemiological study showed that monthly and weekly consumption of red wine is associated with a lower risk of dementia (Daffner 2010). Several studies have shown that caloric restriction enhance healthy living by activating a new class of histone deacetylase enzymes known as sirtuins (Ramesh et al. 2010; Smith et al. 2010; Wang et al. 2010). Resveratrol is a substance found in some plants which may activate sirtuins either directly or by its action on upstream signalling pathways to stimulate caloric restriction and slow down brain atrophy (Albani et al. 2010; Braidy et al. 2012; Massudi et al. 2012). Despite its poor penetration into the brain, some studies have shown that resveratrol can reduce brain  $A\beta$  aggregation (Chandrashekaran et al. 2010; Ladiwala et al. 2010; Sun et al. 2010). An NIA-funded clinical trial examining the importance of resveratrol for the treatment of AD is currently underway.

### 14 Conclusion

AChEIs and the NMDA receptor antagonist memantine are currently approved for the treatment of AD. Despite this, numerous agents have been identified that exert a plethora of neuroprotective effects that may be useful for the treatment of AD. The clinical impact of current agents that may slow down the progression of AD will be significant if these drugs are administered early during disease onset.

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# **Epilepsy**

# Stjepana Kovac and Matthew C. Walker

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#### Abstract

Epilepsy is a chronic disorder defined by recurrent and unprovoked seizures and is one of the commonest disorders of the brain. Despite the burgeoning of new antiepileptic drugs, the burden of medically intractable epilepsy has not reduced significantly. One-third of patients with epilepsy continue to have unprovoked seizures despite advances in treatment. Cell death in seizures has been observed in in vitro and in vivo models of epilepsy and is the hallmark of hippocampal sclerosis, the commonest pathology associated with drug-resistant epilepsy.

Neuroprotection in epilepsy has a long tradition and with the discovery of new pathways involved in seizure induced cell death, strategies have been refined accordingly. In this chapter we will provide a comprehensive overview on the proposed mechanisms underlying cell death in epilepsy and how these can be targeted by exogenous and endogenous neuroprotectants. We will focus on the NMDA receptor and its downstream signalling cascades and their role in seizure induced cell death and potential drug targets arising from this central position within the cascade of cell death. We will summarize evidence

S. Kovac (🖂) • M.C. Walker

UCL Institute of Neurology, University College London, London, UK e-mail: stjepana.kovac.09@ucl.ac.uk; m.walker@ucl.ac.uk

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supporting the involvement of AMPA receptors in seizure induced cell death. In the last part the focus will be on mitochondria and ROS as pivotal players in seizure induced cell death and how these are linked to NMDA receptor dysfunction. We will outline strategies targeting these downstream signalling cascades.

# 1 Introduction

Epilepsy is a chronic disorder defined by recurrent and unprovoked seizures and is one of the commonest disorders of the brain. Despite the burgeoning of new antiepileptic drugs in the twentieth and twenty-first century, the burden of medically intractable epilepsy has not reduced significantly. One-third of patients with epilepsy continue to have unprovoked seizures despite advances in treatment (Duncan et al. 2006). Cell death in seizures has been observed in in vitro and in vivo models of epilepsy and is the hallmark of hippocampal sclerosis, the commonest pathology associated with drug-resistant epilepsy (Swanson 1995; Deshpande et al. 2008). This neuronal death is critically dependent upon excessive neuronal activity and the duration of such activity (Pitkänen et al. 2002). Seizures are usually self-limiting; generalized tonic-clonic seizures typically last less than 90 s (Theodore et al. 1994). However, seizures can be prolonged, and status epilepticus (defined as continuous seizure activity lasting longer than 30 min), the maximal expression of epilepsy, can occur in people with or without a prior history of epilepsy (often as the result of acute brain pathology, e.g., stroke, infection or head injury). This condition is a medical emergency, as it can lead to substantial neuronal cell death, resulting in cognitive and neurological deficits. Even single prolonged seizures in humans can lead to permanent damage in the hippocampus (Pinto et al. 2011). Animal studies support this. The first studies, showing that electrographic seizure discharges alone rather than secondary systemic complications during generalized seizures can lead to cell death, were published by Meldrum and colleagues (1973).

Neuroprotection in epilepsy has a long tradition and was mainly aimed at preventing seizure-induced cell death. However, there is growing evidence that prolonged seizures result in not only neuronal cell death but also neuronal and network dysfunction, which also contribute to the associated morbidities. Therefore, a definition of neuroprotection in epilepsy should include preventing not just neuronal cell death but also network dysfunction (Sutula et al. 2003; Walker 2007). This is difficult to capture in an experimental setting, especially in experimental animal work. In contrast, robust assays are available to measure neuronal cell death in animals. Much research into neuroprotection in epilepsy has therefore focussed on seizure-induced cell death.

This chapter will provide a comprehensive overview on the proposed mechanisms underlying cell death in epilepsy and how these can be targeted by exogenous and endogenous neuroprotectants. The first part will focus on the NMDA receptor and its downstream signalling cascades and their role in seizureinduced cell death and potential drug targets arising from this central position within the cascade of cell death. Evidence supporting the involvement of AMPA receptors in seizure-induced cell death will be summarized. The final part will focus on mitochondria and ROS as pivotal players in seizure-induced cell death and how these are linked to NMDA receptor dysfunction. Strategies targeting these downstream signalling cascades will be outlined.

### 2 Neuronal Cell Death in Epilepsy

For over 100 years, it has been recognized that status epilepticus can result in damage and injury to the hippocampus (Pfleger 1880). Prolonged convulsive seizures can lead to hypotension, ventilatory dysfunction, and metabolic acidosis due to excessive muscle activity that can lead to neuronal damage (Walton 1993). It was thought that this physiological compromise was the main mechanism underlying neuronal death, which would therefore be restricted to convulsive status epilepticus. Pioneering experiments by Meldrum and colleagues in baboons in the 1970s challenged the prevailing orthodoxy; they showed that brain seizure activity can cause neuronal damage even in the absence of a systemic convulsion; seizure-induced brain damage is therefore not just a secondary phenomenon (Meldrum et al. 1973). That seizure activity itself can cause neuronal injury is now widely accepted and it is appreciated that even non-convulsive status epilepticus, a condition encountered in certain clinical settings, such as in comatose patients in the intensive care unit, can result in neuronal damage (Schmutzhard and Pfausler 2011). A more vexed question is whether even single, brief seizures can cause neuronal damage. Longitudinal neuroimaging studies in humans and histopathological studies in animals indicate that repeated seizures over time can result in damage, in particular to the hippocampus (Kälviäinen and Salmenperä 2002; Kotloski et al. 2002).

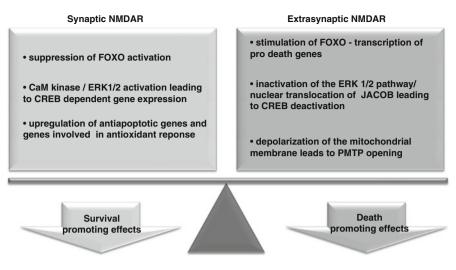
# 3 NMDA Receptors and Seizure-Induced Cell Death

The NMDA receptor has been linked to epilepsy and is an important neuroprotective target in epilepsy. NMDA receptor dysfunction as in NMDA receptor encephalitis, where autoantibodies are directed against the NMDA receptor, can cause seizures (Dalmau et al. 2008). Anti-NMDA receptor antibody mediate reversible decreases in NMDAR surface density and synaptic localization (Hughes et al. 2010), which could explain the associated cognitive deficits seen with this condition but does not easily explain the common occurrence of seizures. NMDAR involvement is seen not only in rare cases of autoantibody-mediated epilepsy but also in more common epilepsies. Epileptogenic lesions such as focal cortical dysplasia have an altered pattern of NMDA receptor expression with an abundance of the NR2B subunit (Ying et al. 2004; Möddel et al. 2005). NMDA receptor activation during prolonged seizures may also regulate the expression and function of other channels. The current first-line treatments for status epilepticus target the benzodiazepine site of the GABA(A) receptor. However, prolonged seizure activity leads to NMDA receptor activation and internalization of GABA(A)

receptors (Muir et al. 2010) and therefore after  $\sim 30$  min of seizure activity benzodiazepines lose their efficacy (Naylor et al. 2005). The reverse holds true for the NMDA receptor. NMDA receptor subunits are mobilized to the synaptic membrane after continuous seizure activity (Chen and Wasterlain 2006).

NMDA receptors have also been identified as the crucial mediator of glutamate-induced neurotoxicity, and it was later discovered that this is mainly mediated by Ca<sup>2+</sup> influx (Lucas and Newhouse 1957; Curtis et al. 1959). NMDA receptor antagonists given early after status epilepticus in animal models have been shown to be neuroprotective (Schauwecker 2010; Loss et al. 2012). There is evidence that NMDA receptor excitotoxicity depends on the type of subunit expressed and on the location relative to the synapse with many proposing that the predominantly extrasynaptic NR2B subunit mediates detrimental effects on neurons (Monver et al. 1994; Tovar and Westbrook 1999; Hardingham et al. 2002; Liu et al. 2007). In keeping with this, ifenprodil, which predominantly targets the NR2B-containing NMDA receptors, has been shown to be neuroprotective if given after status epilepticus (Frasca et al. 2011). However, the NMDA receptor not only has negative effects on neurons, as it is pivotal for long-term potentiation, a fundamental mechanism underlying memory consolidation. Moreover, the NMDA receptor has protective effects (Hardingham and Bading 2010). Such protective effects are mediated by CREB (cyclic-AMP response element-binding protein), a transcription factor which is activated by nuclear Ca<sup>2+</sup>. CREB activation enhances pro-survival gene expression and therefore provides neuroprotection against insults (Chrivia et al. 1993; Hardingham et al. 2001; Papadia et al. 2005). CREB is regulated by the ERK 1/2 pathway and CaM kinase (Wu et al. 2001). Other important pathways that have been implicated in NMDA receptor-mediated neuroprotection include suppression of FOXO and boosting of antioxidant defense genes (Papadia et al. 2008). Interestingly, these pathways are affected by synaptic and extrasynaptic NMDAR activation in opposing fashions, emphasizing the dichotomy between protective synaptic and neurotoxic extrasynaptic NMDAR activation (Hardingham and Bading 2010; Fig. 1). More recently it has been discovered that synaptic NMDARs are gated by different endogenous co-agonists from those that gate non-synaptic NMDARs with the former being gated by *d*-serine and the latter by glycine (Papouin et al. 2012). This differential gating provides novel more specific targets for either the synaptic or extrasynaptic NMDARs. Interestingly, the same group found that NMDA-induced neurotoxicity relies solely on synaptic NMDAR activation, thereby challenging previous reports, which identified the extrasynaptic NMDAR as the major culprit for NMDAR-induced neurotoxicity.

One endogenous substance that acts at the glycine site of the NMDA receptor is kynurenic acid. The kynurenine pathway is the main route of amino acid tryptophan metabolism from dietary intake that is not used for protein metabolism. The two main members along this pathway are quinolinic acid and kynurenic acid. Both are linked to NMDA receptor function. Quinolinic acid acts as an agonist, whereas kynurenic acid acts as an antagonist at NMDA receptors (Stone and Perkins 1981; Vamos et al. 2009). Kynurenic acid acts at both the glutamate and at the allosteric glycine site of the NMDA receptor and at higher concentrations inhibits AMPA receptors



**Fig. 1** Opposing effects of synaptic versus extrasynaptic NMDA receptors. *FOXO* forkhead box O class of transcription factors, *CREB* cyclic-AMP response element-binding protein, *CaM* nuclear Ca<sup>2+</sup>/calmodulin-dependent protein. A more detailed summary of the opposing effects of synaptic versus extrasynaptic NMDA receptors is available in Hardingham and Bading (2010)

(Ganong and Cotman 1986; Watkins et al. 1990). Both receptors have been linked to epilepsy and epileptogenesis; kynurenic acid (KYNA) could therefore be an ideal neuroprotectant in epilepsy. There is evidence that the kynurenine pathway is involved in epilepsy and epileptogenesis. Depletion of KYNA within the hippocampus has been shown in pentylenetetrazole-induced seizures (Maciejak et al. 2009), and endogenously produced KYNA showed strong inhibitory effect on epileptiform activity vitro epilepsy model (Scharfman et al. 1999).

# 4 AMPA Receptors and Seizure-Induced Cell Death

The AMPA receptor is a tetrameric ionotropic transmembrane receptor for glutamate with fast synaptic transmission properties. It is the most abundant ionotropic glutamate receptor in the central nervous system. AMPA receptors are upregulated in neocortical tissue from patients suffering from pharmacoresistant epilepsy (Zilles et al. 1999; Palomero-Gallagher et al. 2012). Ion permeability of the receptor is determined by its subunit composition; GluR2-lacking receptors are permeable to  $Ca^{2+}$ , Na<sup>+</sup>, and K<sup>+</sup> and are inwardly rectifying, while channels containing the GluR2 subunit are selective for Na<sup>+</sup> and K<sup>+</sup> only (Isaac et al. 2007). Such differences in permeability are important, since in refractory status epilepticus, GluR2-subunits are internalized and the GluR2-lacking AMPARs are preferentially expressed on the surface of the cell. Furthermore, Rajasekaran and colleagues have shown that these receptor changes in the hippocampi of rats subjected to status epilepticus translate into changes in the functional properties of AMPAR-mediated currents, which show

increased inward rectification and the activation of which results in intracellular accumulation of Ca<sup>2+</sup> (Rajasekaran et al. 2012). Drugs that target AMPA receptors have consequently also been shown to be neuroprotective in status epilepticus (Fritsch et al. 2010). AMPA receptor changes have not only been found in acquired epilepsy models but also in genetic models of epilepsy. The stargazer mice (lacking stargazing) have a trafficking defect in synaptic AMPARs in reticular thalamic nucleus cells and enhanced synaptic NMDARs activity leading to increases in thalamic excitability (Lacey et al. 2012). Targeting AMPA receptor trafficking therefore may reduce seizure-induced cell death.

# 5 Seizure-Induced Cell Death, Epilepsy, and Mitochondria

The common feature of both NMDA and GluR2-lacking AMPAR activation is an intracellular increase in Ca<sup>2+</sup>. Ca<sup>2+</sup> homeostasis is closely linked to the mitochondria and mitochondrial dysfunction leads to epilepsy in humans. It is well established that mutations in the mitochondrial genome or mutations in the nuclear genes coding for mitochondrial proteins lead to epileptic phenotypes (DiMauro 2004; Schapira 2006). Furthermore, a tight link between mitochondrial ATP production and seizures can be postulated, since seizures are a prominent feature in Leigh syndrome patients showing mutations in ATPase 6 gene (Canafoglia et al. 2001). Classical examples of mutations in the mitochondrial genome leading to seizures are MELAS (mitochondrial myopathy, encephalopathy, lactic acidosis, and stroke-like episodes) and MERRF (myoclonic epilepsy with ragged red fibres) syndrome (Rahman 2012). As implied in the names of mitochondrial disorders, multiple systems are affected; cerebral involvement represents only one facet of these multiorgan conditions. Epilepsies due to mutations in mitochondrial genes or nuclear genes coding for mitochondrial function are rare, when compared to the vast majority of epilepsies in general neurological practice. However, there is growing evidence that mitochondria and mitochondrial dysfunction may play an important role in many other epilepsies (Kann and Kovács 2007; Kann et al. 2011). Ongoing seizure activity induces ionic shifts and thereby increasing the demand for energy substrates to maintain transmembrane ion homeostasis (Lux et al. 1986). Therefore, it seems reasonable to link seizure activity to mitochondria, which are the main organelles responsible for the maintenance of transmembrane ion homeostasis. That mitochondria may play a prominent role in seizure generation is supported by the observation that mitochondrial membrane potential and  $Ca^{2+}$ concentrations change dramatically during epileptiform activity in vitro (Kovacs et al. 2005; Kovac et al. 2012). Mitochondrial complex I deficiency has been found in tissue resected from patient undergoing epilepsy surgery thus linking mitochondrial respiration to chronic seizure activity (Kunz et al. 2000). Another link comes from the effective treatment of refractory seizures with the ketogenic diet, a highfat, low-carbohydrate diet (Bough et al. 2006; Bough 2008). Further analysis of the changes in gene expression secondary to the ketogenic diet showed mainly upregulation of genes involved in oxidative phosphorylation and TCA cycle function (Bough et al. 2006). Finally, the pathology underlying one of the most common forms of epilepsy is hippocampal sclerosis which is characterized by apoptosis and necrosis, features that are tightly linked to mitochondria (Sloviter 1996).

But how do mitochondria contribute to seizure-induced cell death? There are two recognized forms of cell death: apoptosis and necrosis, and the mitochondria are important players in both. Apoptosis is defined as programmed cell death. A cascade of events leads to characteristic morphological cell changes with shrinkage of the cell with an intact cell membrane, fragmentation of DNA, and finally phagocytosis leading to cell death. In contrast necrotic cell death is described as a premature uncontrolled neuronal death that involves activation of hydrolytic enzymes and a rupture of the cell membrane followed by an inflammatory response.

Apoptosis involves intrinsic (activation within the cell or mitochondrion) and extrinsic (activation through plasma membrane receptors) pathways that converge to activate the executioner caspases, caspase 3 and caspase 7 (Tait and Green 2010). Inhibition of caspase 3 has been shown to be neuroprotective even if treatment is delayed for after the status epilepticus (Narkilahti et al. 2003b). Both pathways rely on mitochondrial outer membrane permeabilization (MOMP), which indicates a key point of no return. MOMP leads to release of cytochrome C. Cytochrome C is involved in electron transport during oxidative phosphorylation. However, it also leads to caspase activation after permeabilization of the outer mitochondrial membrane. Cells lacking cytochrome C fail to activate caspases and exhibit resistance to intrinsic apoptosis (Li et al. 2000).

Mitochondria act as a gatekeeper tuning  $Ca^{2+}$  concentrations in the cell and in cytoplasmic domains by buffering the  $Ca^{2+}$  during  $Ca^{2+}$  signalling. However, this buffering can be a starting point for apoptotic and necrotic cell death, if the matrix Ca<sup>2+</sup> exceeds mitochondrial Ca<sup>2+</sup> buffering capacity. This process leads to permeability transition, which is a phenomenon induced by high levels of matrix Ca<sup>2+</sup>. It is characterized by the opening of the permeability transition pore (MPTP). The characteristic feature of MPTP opening is a permeability increase of the inner mitochondrial membrane to substances with small molecular masses (up to 1,500 Da) (Bernardi 1999; Crompton et al. 1999; Bernardi et al. 2006). Factors that have been indentified to contribute to mitochondrial permeability transition are Ca<sup>2+</sup> accumulation and mitochondrial membrane potential depolarization, whereas a low pH, Mg<sup>2+</sup>, and ADP prevent MPTP opening. The exact structure of the MPTP still needs to be characterized; however, it is most likely a multimeric complex, which may be assembled following activation (Zorov et al. 2009). Cyclophilin D (CypD) is a component of the MPTP, which is used pharmacologically, as the CypD inhibitors, such as cyclosporin A, are potent inhibitors of MPTP opening (Elrod et al. 2010). Strong evidence that MPTP could be a target for drug development comes from recent studies in ischemic heart disease. Inhibition of MPTP opening has been hypothesized to underlie the protective effect of ischemic preconditioning and post conditioning of myocytes (Hausenloy et al. 2009).

These strategies have been translated to the brain and MPTP opening has become a target in stroke and traumatic brain injury (Sullivan et al. 2005; Osman et al. 2011). Cyclosporin A prevents seizure-induced mitochondrial membrane depolarization, an event that leads to seizure-induced cell death (Kovac et al. 2012). Persistent MPTP opening leads to a depolarization of the mitochondrial membrane potential with a subsequent decrease in ATP production and matrix swelling induces breaks in the outer mitochondrial membrane. Release of cytochrome C which contributes to the formation of the apoptosome finally activates the executor caspases. Besides permanent opening, the MPTP can undergo transient "flickering" states which may prevent permanent opening through decreasing intra-matrix Ca<sup>2+</sup> concentrations (Petronilli et al. 1999). Whether the cell will undergo apoptotic or necrotic cell death depends on the proportion of mitochondria undergoing MPTP opening. Mitochondria undergoing MPTP opening are not capable of ATP generation due to a dissipation of the mitochondrial membrane potential. However, ATP is required to aggregate the apoptosome which is necessary to induce programmed cell death via caspases. If not enough ATP is available for apoptosome formation, the rise in cytosolic  $Ca^{2+}$  will activate hydrolytic enzymes leading to necrosis (Gramaglia et al. 2004). It is important to note that increases in reactive oxygen species (ROS) contribute to MPTP opening by increasing Ca<sup>2+</sup> load via stimulation of IP<sub>3</sub>/ryanodine receptors and via inhibition of SERCA pumps and plasma membrane  $Ca^{2+}$  channels (PMCA) (Camello-Almaraz et al. 2006).

Inhibition of cell death pathways would seem a rational approach to prevent seizure-induced cell death. However, studies using this strategy have varying and inconsistent results. Inhibiting caspase 3 activation, for example, has been shown to be neuroprotective after status epilepticus in one animal model, whereas another study showed that this pathway is not involved in seizure-induced neuronal necrosis (Fujikawa et al. 2002; Narkilahti et al. 2003a, b; Walker 2007). Possible explanations for this variation are differences in the epilepsy models used in these studies and the time point at which the intervention takes place. Indeed, there may be more than one downstream pathway involved in mediating neuronal death in epilepsy and the relative importance of these pathways may depend upon a host of factors, highlighting the significant challenges facing the search for neuroprotective targets in epilepsy.

Given that energy failure and starvation play a pivotal role in seizure-induced cell death, anaplerosis seems to be an appealing strategy to reduce seizure-induced cell death. Anaplerosis is the replenishment of energetic substrates and is a strategy that has been used to prevent seizure-induced cell death and seizures themselves (for a comprehensive review see (Kovac et al. 2012)). Anaplerotic substrates, such as pyruvate or alpha-ketoglutarate, for example, reduce seizure-induced cell death both in vitro and in vivo (Yamamoto 1990; Yamamoto and Mohanan 2003; Kim et al. 2007; Carvalho et al. 2011; Kovac et al. 2012).

Factors that limit the use of anaplerotic substrates for the treatment of seizures and seizure-induced cell death are pharmacokinetic properties of these compounds and their bioavailability. Some of the compounds of the tricarboxylic acid cycle are not membrane permeable (e.g., succinyl-CoA) or lack adequate transporters in the neuron, (e.g., citrate), preventing their use as an anaplerotic substrate (Gnoni et al. 2009).

Pyruvate seems to be the most promising anaplerotic substrate given its pharmacokinetic properties and evidence from in vivo and in vitro animal and human data. In addition pyruvate exhibits antioxidant properties as it acts as a free radical scavenger (Andrae et al. 1985; Bassenge et al. 2000; Varma and Hegde 2007).

# 6 Seizure-Induced Cell Death and Reactive Oxygen Species (ROS)

ROS can contribute to neuronal damage and therefore to seizure-induced cell death through a myriad of pathways. ROS, especially of mitochondrial origin, contribute to MPTP opening by increasing Ca<sup>2+</sup> load via stimulating IP<sub>3</sub>/ryanodine receptors and inhibiting SERCA pumps and plasma membrane Ca<sup>2+</sup> channels (PMCA) (Camello-Almaraz et al. 2006). More recently ROS generated as a direct consequence of NMDA receptor activation have been described (Brennan et al. 2009).

Due to their reactive nature, ROS can contribute to cell death in a more direct manner. These are lipid peroxidation, protein denaturation inducing inactivation of enzymes, and nucleic acid damage. It needs to be highlighted that all these events are consequences of a very high intracellular ROS burden, exceeding the cell's capacity for antioxidant defense.

With regard to the brain, lipid peroxidation is probably the most important detrimental ROS effect, due to the abundance of lipids within the brain. Lipid peroxidation leads to the production of conjugated dienic hydroperoxides that decompose into aldehydes, dienals, or alkanes. These are toxic to the neuron and can lead to apoptosis (McCracken et al. 2000). Nuclear damage to DNA can lead to activation of repair enzymes such as poly (ADP-ribose) polymerase (PARP). This enzyme is known to lead to rapid ATP depletion which can stimulate cascades leading to cell death (Szabó 1998; Heeres and Hergenrother 2007). On the other hand, ROS induce release of cytochrome C from the mitochondria which activates the apoptosome inducing the caspase cascade, which leads to caspase 3-induced cleavage of nuclear DNA repair enzyme (Kim et al. 2000).

Oxidative stress has been implicated in the pathology associated with acute seizures, status epilepticus, and epilepsy. Acute increases in reactive oxygen species leading to damage of proteins, lipids, and ultimately cell death were shown after repeated seizures (Bruce and Baudry 1995; Liang et al. 2000; Tejada et al. 2007). It has been shown that antioxidant compounds such as vitamin C prevent seizure-induced neuronal death (Santos et al. 2008, 2009). Moreover, the lack of cellular antioxidants, as in superoxide dismutase mutant mice, leads to seizures in vivo (Lynn et al. 2005). In humans, magnetic resonance spectroscopy has shown decreased brain levels of glutathione, indicative of oxidative stress, in the brain of patients with epilepsy when compared to age-matched healthy controls (Mueller et al. 2001). Reactive oxygen species can damage lipids in cell membranes.

ROS degrade polyunsaturated lipids, forming malondialdehyde (MDA), which therefore is a marker of oxidative damage within a cell. MDA has been shown to increase following chronic epilepsy implying that seizure-induced ROS production leads to oxidative damage of lipids and cell membranes (Bruce and Baudry 1995; Frantseva et al. 2000; Tejada et al. 2007). Animal studies have shown decreased levels of glutathione in rats after prolonged seizures (Sleven et al. 2006), and the ketogenic diet, an effective treatment for pharmacoresistant seizures, increased glutathione levels in rat hippocampi (Jarrett et al. 2008).

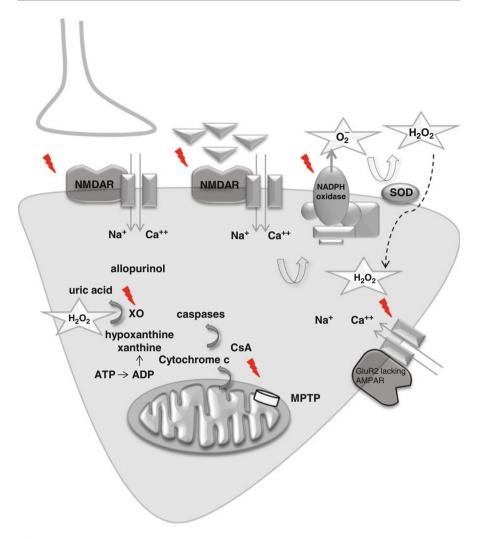
Two mechanistically different therapeutic approaches can reduce the oxidative burden within the cell. One is blocking the source of free radical generation, whereas the other aims at scavenging the free radicals that have already been produced within the cell. The first approach requires identifying the source of free radical generation within the cell.

Several enzymes generate ROS within the cell. The candidate enzymes for ROS generation during seizures are the mitochondria, NADPH oxidase, and xanthine oxidase. Mitochondria, through reactions in the electron transport chain at complex I and III, are the main producers of ROS within the cell. That the NADPH oxidase may play an important role in seizure-induced ROS generation is suggested by the finding that NMDA receptors play a pivotal role in seizure-induced neuronal damage (Schauwecker 2010) and that recent evidence indicates that direct pharmacological activation of NMDA receptors can activate NADPH oxidase, increasing free radical production and consequently neuronal death (Abramov et al. 2007; Brennan et al. 2009; Girouard et al. 2009). Moreover, xanthine oxidase (XO) is also a major potential source of ROS during periods of increased metabolism such as occur during seizure activity.

Excessive NMDA receptor activation is a crucial event in acute neurological disease with detrimental downstream effects on neurons which include free radical generation (Lipton 1999) and there is a strong link between epilepsy and NMDA receptor dysfunction (Ying et al. 1999; Schauwecker 2010). Recent data suggest that NADPH oxidase is the primary source of superoxide induced by NMDA receptor activation (Brennan et al. 2009), and NADPH oxidase has been identified as a pivotal source of ROS generation in acute neurological disease such as stroke (Abramov et al. 2007).

In keeping with these findings, increases in NADPH oxidase activity have been linked to epilepsy models such as the pilocarpine and kainite model of epilepsy (Patel et al. 2005; Di Maio et al. 2011), and inhibition of NADPH oxidase was effective in reducing cell death in the in vivo pilocarpine model of temporal lobe epilepsy (Pestana et al. 2010).

ATP depletion results in adenine formation and consequently an increase in hypoxanthine and xanthine, substrates for xanthine oxidase (Xia and Zweier 1995; Kinugasa et al. 2003). Previous reports have suggested that there may be a beneficial effect of xanthine oxidase inhibition on seizures (Kramer et al. 1990; Tada et al. 1991; Zagnoni et al. 1994), although such an effect is not universally observed.



**Fig. 2** Possible targets for neuroprotection in epilepsy. The figure illustrates potential targets of neuroprotection in epilepsy which include the ionotropic NMDA and AMPA glutamate receptor, the mitochondria and downstream mechanisms of ROS generation as the NADPH or xanthine oxidase. Abbreviations: *NADPH OX* NADPH oxidase, *MPTP* mitochondrial permeability transition pore, *CsA* cyclosporin A, *NMDAR* NMDA receptor, *AMPAR* AMPA receptor, *XO* xanthine oxidase, *SOD* superoxide dismutase

This chapter has outlined several mechanisms involved in seizure-induced cell death. The evidence for targeting NMDA and AMPA receptor, mitochondrial function, and ROS including several downstream targets of those receptors has been discussed. Neuroprotective targets in epilepsy are summarized in Fig. 2.

# 7 Conclusion

There are several interacting yet distinct mechanisms that can contribute to neuronal death in epilepsy, making common neuroprotective strategies challenging. The relative roles of each of the above mechanisms also show time dependency in status epilepticus with critical early steps likely to be the activation of ionotropic receptors and calcium influx, while later steps involve the activation of signalling and metabolic pathways. Directing treatments at the early steps has the advantage that there may be fewer targets but this strategy may be temporally less practical. Treating the later stages has a temporal advantage but the large number of pathways possibly involved presents a different challenge.

The observation that repeated brief seizures over time can lead to neuronal damage suggests that long-term neuroprotective strategies could offer the possibility of modifying comorbid declines in memory and other associated conditions such as depression.

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# Excitotoxicity and Amyotrophic Lateral Sclerosis

# Kim A. Staats and Ludo Van Den Bosch

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#### Abstract

Amyotrophic lateral sclerosis (ALS) is a devastating neurodegenerative disorder characterized by the progressive and selective death of motor neurons in the motor cortex, brainstem, and spinal cord. The only therapy with proven efficacy for ALS patients is a treatment with riluzole that offers patients a moderate

K.A. Staats (🖂) • L. Van Den Bosch

Laboratory of Neurobiology and Leuven Research Institute of Neurodegenerative Diseases (LIND), KU Leuven, Leuven, Belgium

Vesalius Research Center, VIB, Leuven, Belgium e-mail: kim.staats@vib-kuleuven.be; Ludo.VandenBosch@vib-kuleuven.be

increase in survival. Riluzole interferes with excitotoxicity, which is neuronal death caused by overstimulation of glutamate transporters. The high sensitivity of motor neurons to excitotoxicity is further illustrated by the observation that a number of exogenous toxic substances (excitotoxins) that stimulate glutamate receptors result in selective motor neuron death. This selective vulnerability of motor neurons to excitotoxic insults is due to their poor calcium buffering capacity in combination with the expression of calcium-permeable AMPA type of glutamate receptors. The AMPA receptor subunit responsible for calcium impermeability, GluR2, is expressed at lower levels, and RNA editing of GluR2 required for calcium impermeability is diminished in motor neurons of ALS patients. Not only calcium permeability of AMPA receptors but also glutamate transport into neighboring astrocytes plays an important role in ALS. Decreased levels of glutamate transporters are present in ALS patients and ALS animal models, implying an increased vulnerability to glutamate. This chapter aims to provide an overview of the known players and their interactions in the field of excitotoxic motor neuron death and in the context of ALS.

#### Keywords

ALS • Excitotoxicity • Motor neuron loss • Motor neuron disease • Glutamate

# 1 Introduction of ALS

#### 1.1 Basics

Amyotrophic lateral sclerosis (ALS) is a devastating fatal neurodegenerative disease, with progressive loss of motor neurons of the motor cortex, brainstem, and spinal cord and denervation of muscle fibers, resulting in increasing muscle weakness and paralysis. The disease has an incidence of 2.7 cases per 100,000 people in Europe (Logroscino et al. 2010) and is more prevalent in the later years of life. In lack of a medical cure, average life expectancy is between 2 and 5 years. Patients usually die due to respiratory insufficiency or may opt for euthanasia where legislature permits (Maessen et al. 2010). Although ALS is characterized by degeneration of central nervous system tissue, mental functions remain largely unaffected resulting in a locked-in state (Kotchoubey et al. 2003). Only one drug is currently available to treat the disease, riluzole, and slows disease progression moderately (Miller et al. 2007).

#### 1.2 ALS Genetics

There are both familial and sporadic cases of the disease, and a common disease mechanism is expected due to the indistinguishable clinical phenotypes of these patients. Approximately 90 % of ALS cases are sporadic in which a gene-environment

interaction is hypothesized. The remaining 10 % of ALS cases are inherited by mutations in a number of genes including superoxide dismutase 1 (SOD1), vesicle-associated membrane protein-associated protein B (VAPB), TAR DNA binding protein (TDP43), fused in sarcoma/translocated in liposarcoma (FUS/TLS), optineurin (OPTN), valosin-containing protein (VCP), ubiquilin2 (UBQLN2), and p62 (SQSTM1) which have been identified as causes of familial ALS (Van Deerlin et al. 2008; Rutherford et al. 2008; Del Bo et al. 2009; Kwiatkowski et al. 2009; Vance et al. 2009; Maruyama et al. 2010; Fecto et al. 2011; Deng et al. 2011; Johnson et al. 2010). In addition, hexanucleotide repeat expansions in the noncoding part of the "chromosome 9 open reading frame 72" (C9ORF72) gene were discovered as a new cause of ALS (Dejesus-Hernandez et al. 2011; Renton et al. 2011).

# 1.3 ALS Models

Overexpression of mutant forms of human SOD1 causes the ALS phenotype of transgenic SOD1 mice, which have made a large contribution to ALS research (Gurney et al. 1994). Many hallmarks of the disease are shared by patients and this rodent model, including specific motor neuron loss, aggregate formation, astrogliosis, microgliosis, and progressive paralysis. SOD1 detoxifies potentially cell-damaging free radicals, but the genetic ablation of SOD1 does not produce an ALS-like phenotype in mice (Reaume et al. 1996; Shefner et al. 1999), implying that mutant SOD1 causes the disease by a toxic gain of function. This gain of function may exert itself by protein misfolding, aggregation, impaired proteasome functioning, impaired retrograde transport, excitotoxic cell death, and others (reviewed in Bruijn et al. (2004)). Unfortunately, the discovery of the other ALS genes described above has not yet progressed into useful ALS model organisms, forcing most work described below to have been conducted on mutant SOD1 cells, on rodents, or on patients.

#### 1.4 Non-cell Autonomous

ALS is a non-cell autonomous disease (Boillee et al. 2006a), meaning that multiple cell types affect disease pathology. By addition or deletion of mutant SOD1 in specific cell types, a number of cell types have been identified that influence the disease, including astrocytes (Yamanaka et al. 2008), microglia (Boillee et al. 2006b), Schwann cells (Lobsiger et al. 2009), and motor neurons (Jaarsma et al. 2008). Despite the influence of multiple cell types, mutant SOD1 expressed solely in neurons is sufficient to initiate the disease, with a slowed disease progression (Jaarsma et al. 2008). The vulnerability of motor neurons is also portrayed by the selective loss of motor neurons in the motor cortex, brainstem, and spinal cord in patients. There are a number of hypotheses that explain this cell type selectivity, including the long axons of these cells and the poor buffering of intracellular calcium.

# 1.5 Excitotoxicity in ALS

Glutamate is the neurotransmitter used to excite motor neurons that is the initiator of excitotoxicity in ALS. This neurotransmitter is the most abundant excitatory neurotransmitter in the brain and binds to *N*-methyl-D-aspartate (NMDA) or  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA)/kainate (ionotropic) receptors and metabotropic glutamate receptors (mGluR). The levels of glutamate are increased in ALS patients (Fiszman et al. 2010; Spreux-Varoquaux et al. 2002), implying glutamate's role in the disease. The importance of excitotoxicity in ALS has been demonstrated mainly by the beneficial effects obtained by treating patients with riluzole. Although the precise mechanisms of this drug are not yet known, it has been shown to block NMDA receptors, it enhances reuptake of glutamate from the synaptic cleft, and it inhibits glutamate release by blocking voltage-gated sodium channels (Siniscalchi et al. 1999), thus preventing motor neuron cell death. Riluzole treatment increases predicted life span by 12 % in ALS mice (Lacomblez et al. 1996; Bensimon et al. 1994) and increases the probability of 1-year survival in patients by 9 % (Miller et al. 2007).

# 1.6 Non-glutamate Excitotoxins

A small number of substances are excitotoxic and can cause motor neuropathies and/or neurodegeneration. Domoic acid causes a motor neuropathy in some patients after food poisoning due to consumption of mussels containing high levels of this compound (Debonnel et al. 1989). Domoic acid is also associated with epileptic seizures in humans (Teitelbaum et al. 1990), sea lions (Scholin et al. 2000), and rats (Muha and Ramsdell 2011) and lowers the threshold for seizures in adult rats when exposed to it at neonatal stages (Gill et al. 2010). Also β-N-oxalyl-amino-L-alanine (BOAA), also known as  $\beta$ -oxalyl-L- $\alpha$ , $\beta$ -diaminopropionic acid (ODAP), causes excitotoxicity. BOAA is highly present in the chickling pea (Lathyrus sativus) of which consumption leads to lathyrism (Streifler and Cohn 1981; Spencer et al. 1986). This substance is an AMPA receptor agonist (Willis et al. 1994; Kunig et al. 1994) that causes motor neuron degeneration (Chase et al. 1985; Weiss et al. 1989) and alternative effects on the central nervous system (Willis et al. 1994; Kunig et al. 1994). Another excitotoxin is one discovered by a high prevalence of ALS on the island of Guam, also known as the Western Pacific amyotrophic lateral sclerosis-Parkinsonism-dementia or the amyotrophic lateral sclerosis-Parkinsonismdementia complex (ALS/PDC). Consumption of cycad seed products (Cycas circinalis) is responsible for the neurodegeneration in ALS/PDC of which β-methylamino-alanine (BMAA) is identified as a damaging substance (Steele and Guzman 1987; Hudson 1991; Wilson et al. 2002). BMAA is also a glutamate receptor agonist that causes motor neuron syndrome (Murch et al. 2004; Chang et al. 1993; Purdie et al. 2009) and damages motor neurons (Rao et al. 2006), cholinergic neurons in vitro (Liu et al. 2010; Brownson et al. 2002), and hippocampal neurons in vivo (Buenz and Howe 2007).

# 1.7 Poor Calcium Buffering

An additional explanation for the selective motor neuron vulnerability to excitotoxicity is poor capability of this cell type to buffer intracellular calcium; ALS-vulnerable spinal cord and brainstem motor neurons in mice display low endogenous calcium buffering capacities (Palecek et al. 1999; Lips and Keller 1998). In addition, ALS-resistant oculomotor neurons contain a far larger calcium buffering capacity than ALS-vulnerable motor neurons (von Lewinski and Keller 2005). This difference in calcium buffering capacity is directly due to the differential expression levels of the calcium-binding proteins calbindin-D28k and parvalbumin (Obal et al. 2006). The beneficial effect of calcium buffering by calcium-binding proteins is confirmed in experiments of acute neurodegeneration in a transgenic mouse by overexpressing the calcium-binding protein parvalbumin. According to expectations, parvalbumin overexpressing cells are protected against neurodegeneration induced by axotomy in vivo (Paizs et al. 2010) and excitotoxic cell death in vitro (Van Den Bosch et al. 2002a). In addition, parvalbumin overexpression delays disease onset and extends survival in ALS mice (Beers et al. 2001).

# 2 AMPA Receptors in Excitotoxicity in ALS

#### 2.1 Basic Excitotoxicity

After release from the presynaptic neuron into the synaptic cleft, glutamate binds to NMDA, AMPA, or metabotropic receptors. The AMPA receptor plays an important role in motor neuron death in ALS. The AMPA receptor is a tetramer combining four different subunits (glutamate receptor units 1–4 (GluR1–4)) (Shi et al. 1999). All subunits bind glutamate, and the channels open after occupation of at least 2 binding sites (Mayer 2005). The importance of this receptor in ALS is demonstrated by the ablation of glutamate-induced apoptosis in cortical neurons in vitro (Cid et al. 2003) and in vivo when administering an AMPA receptor antagonist (Van Damme et al. 2003). The detrimental role of glutamate in the disease is demonstrated by the pronounced cell death that occurs to primary cultured neurons in vitro when exposed to low levels of glutamate, even as low as physiologically detected in cerebrospinal fluid (CSF) (Cid et al. 2003). To further illustrate the detrimental role of glutamate in ALS, administration of compounds that block the formation of glutamate increases cell survival, both in vivo and in vitro (Cid et al. 2003). In addition, decreasing vesicular glutamate transporter 2 (VGLUT2) extends survival of motor neurons in ALS mice but does not extend life span of ALS mice (Wootz et al. 2010).

#### 2.2 AMPA Receptor-Mediated Cell Death

As discussed previously, motor neurons are excited by glutamate by, among others, the AMPA receptors. Interestingly, the AMPA receptors on motor neurons are mainly calcium permeable in vitro (Van Den Bosch et al. 2000; Carriedo et al. 1996; Van Den Bosch et al. 2002b), which may explain the selective vulnerability of motor neurons to excitotoxic cell death. In addition, extracellular calcium entry via these calcium-permeable AMPA receptors is responsible for selective motor neuron death, as motor neuron death is inhibited by selective blockers of calcium-permeable AMPA receptors (Van Den Bosch et al. 2000). AMPA receptors of motor neurons have a lower rectification index and a higher relative calcium permeability ratio than other neurons (Van Damme et al. 2002). Not only in vitro but also in vivo motor neurons express calcium-permeable AMPA receptors (Greig et al. 2000). Interestingly, infusing the spinal cord of rats with AMPA receptor agonists induces neurodegeneration of a selective blocker of calcium-permeable AMPA receptors (Corona and Tapia 2007; Sun et al. 2006). This demonstrates a specific vulnerability of motor neurons to AMPA receptor-mediated cell death.

#### 2.3 GluR2 Subunit of the AMPA Receptor

The AMPA receptor plays an imperative role in excitotoxicity by its calcium permeability that is determined by the incorporation of the GluR2 subunit in the receptor complex. In most conditions, the AMPA receptor complex contains at least one GluR2 subunit and this prevents the influx of extracellular calcium into the neuron (Seeburg et al. 2001). In contrast, receptors lacking the GluR2 subunit are highly calcium permeable (Seeburg et al. 2001). Murine motor neurons contain lower levels of GluR2 mRNA compared to other neurons (Van Damme et al. 2002). In addition, laser capture microscopy and quantitative PCR demonstrated that the expression level of the GluR2 subunit is lower in (human) spinal motor neurons compared to other neurons (Heath et al. 2002; Kawahara et al. 2003), implying the intrinsic vulnerability of motor neurons to excitotoxicity. A general decrease of GluR2 is found in ALS model mice, portraying an increased vulnerability of these mice to excitotoxic insults (Zhao et al. 2008; Tortarolo et al. 2006). The role of GluR2 in ALS is investigated by genetically ablating GluR2 in ALS mice, which decreases survival in vivo and decreases cell survival in vitro (Van Damme et al. 2005). Upregulation of GluR2 expression in motor neurons of ALS mice increases survival (Tateno et al. 2004). In addition, pharmacological inhibition of the AMPA receptor prolongs survival in ALS mice (Van Damme et al. 2003; Tortarolo et al. 2006; Canton et al. 2001). Interestingly, the surrounding astrocytes influence the expression level of the GluR2 subunit in motor neurons, as soluble factors released from astrocytes affect GluR2 gene expression and neuronal vulnerability to excitotoxic insults, both in vitro and in vivo (Van Damme et al. 2007). Intriguingly, the presence of mutant SOD1 interferes with the production and/or secretion of these factors (Van Damme et al. 2007). Although not all these influencing factors are known, vascular endothelial growth factor (VEGF), brain-derived neurotrophic factor (BDNF), and glial cell-derived neurotrophic factor (GDNF) are able to upregulate GluR2 expression (Bogaert et al. 2010; Brene et al. 2000). Although the role of AMPA receptors is established, there is no genetic evidence that polymorphisms in the GluR2 gene increase susceptibility for ALS (Bogaert et al. 2012).

# 2.4 GluR2 Editing

The GluR2 subunit's special characteristic of preventing calcium influx through the AMPA receptor is due to RNA editing at the Q/R site, and this process is also associated to ALS. This editing in the GluR2 pre-mRNA introduces a positively charged arginine at the Q/R site of the GluR2 peptide instead of the genetically encoded neutral glutamine (Burnashev et al. 1992). Under normal conditions, GluR2 pre-mRNA editing is virtually complete, but under pathological conditions it could become less efficient resulting in more calcium-permeable AMPA receptors. Indeed, it is reported that editing of pre-mRNA of GluR2 is defective in the spinal motor neurons of sALS patients (Kawahara et al. 2004). In addition, GluR2 gene expression and GluR2 RNA editing are reduced in the spinal cord of ALS patients (Takuma et al. 1999). Furthermore, the enzyme responsible for the RNA editing, adenosine deaminase acting on RNA 2 (ADAR2), is only present in 50 % of the motor neurons from sporadic ALS patients (Aizawa et al. 2010). Reduced editing at the Q/R site of GluR2 in mice results in a lethal phenotype accompanied with seizures and neurodegeneration (Brusa et al. 1995; Higuchi et al. 2000). Transgenic mice carrying a minigene with the GluR2 gene encoding an asparagine (GluR2-N) at the Q/R site, which makes editing impossible, are viable and fertile (Kuner et al. 2005) and are a useful tool to investigate the effect of calcium-permeable AMPA receptors, as AMPA receptors incorporating GluR2-N are permeable to calcium (Burnashev et al. 1992). The combined expression of the GluR2-N transgene and endogenous GluR2 alleles results in a twofold increase in permeability for calcium (Feldmeyer et al. 1999). Interestingly, these transgenic mice develop motor neuron degeneration late in life (Feldmeyer et al. 1999), and GluR2-N overexpression induces a progressive decline in function, as well as a degeneration of spinal motor neurons (Kuner et al. 2005). In accordance, GluR2-N expression in ALS mice exacerbates disease progression and reduces survival confirming the role for edited GluR2 in ALS (Kuner et al. 2005). In addition, conditional knockdown of ADAR2 in murine motor neurons causes a late-onset neurodegenerative condition (Hideyama et al. 2010).

# **3 Glutamate Transporters in ALS**

#### 3.1 Glutamate Transporter Levels in ALS

After release from the presynaptic neuron, glutamate is recycled by the glial and endothelial cells, including astrocytes. Astrocytic glutamate reuptake occurs by the glutamate transporters excitatory amino acid transporter 1 (EAAT1) and excitatory amino acid transporter 2 (EAAT2; also known as glutamate aspartate transporter

(GLAST1) and glutamate transporter 1 (GLT-1), respectively). These transporters internalize glutamate, for conversion to glutamine that is returned to the presynaptic neuron (Laake et al. 1995). Reduced glutamate uptake and EAAT2 protein expression are common in both ALS patients and ALS model systems (Staats and Van Den Bosch 2009). In vitro transfection of astrocytes with either mutant SOD1 or wild-type human SOD1 downregulates EAAT2 posttranscriptionally (Tortarolo et al. 2004), though it is unknown whether the downregulation of EAAT2 affects neuronal survival. Mutant SOD1 transfection also decreases glutamate transport in cell lines (Sala et al. 2005). Interestingly, this downregulation is present in ALS model rats at presymptomatic stages through to end stage (Howland et al. 2002), at end stage only (Warita et al. 2002), in ALS model mice at end stage (Bendotti et al. 2001), and in post mortem patient spinal cords by staining for EAAT2 (Sasaki et al. 2001). Interestingly, a decrease of EAAT2 protein is present in mutant SOD1 mice and in an environmental model of ALS/PDC (mice are fed with washed cycad flour containing BMAA, which causes an ALS-like phenotype (Wilson et al. 2002)) (Wilson et al. 2003). Although the loss of EAAT2 in ALS is apparent, it remains unclear whether this loss of EAAT2 proceeds or follows the loss of motor neurons.

# 3.2 Reducing Glutamate Transporters in ALS

To assess whether the loss of glutamate transport or the loss of EAAT2 specifically results in motor neuron loss, pharmacological and genetic tools have been employed. To begin, research conducted by pharmacologically inhibiting glutamate transport in the rat spinal cord failed to show any motor neuron loss despite the increased levels of glutamate (Tovar et al. 2009). This experiment has not (yet) been performed in ALS mice, leaving it to be elucidated whether decreased glutamate transport pharmacologically is detrimental for motor neuron survival in ALS. In contrast, a similar experiment has been performed to address whether EAAT2 loss specifically would induce motor neuron loss. EAAT2 null mice have been generated that live for approximately 6 weeks before they succumb to epileptic seizures and show increased vulnerability to acute brain injury (Tanaka et al. 1997). The heterozygous mice (containing 1 allele of EAAT2) were used to assess the effect of approximately 40 % knockdown of EAAT2 in the spinal cord in ALS mice (Pardo et al. 2006). This described knockdown resulted in a nonsignificant decrease of symptom onset and in significant, but moderate, decrease of life span in ALS mice (Pardo et al. 2006).

# 3.3 Increasing Glutamate Transporters in ALS

To assess the expected beneficial role of EAAT2 in ALS, transgenic mice overexpressing human EAAT2 in astrocytes only were crossbred with mutant SOD1 mice. Although glutamate uptake is increased in the mice and is protective on primary cortical neurons in culture, these effects were unable to affect

symptom onset or life span (Guo et al. 2003). Possibly, the expression levels were too low to induce an effect, or human EAAT2 is not as efficient as murine EAAT2 in mouse, as administration of ceftriaxone (a β-lactam antibiotic) or GPI-1046 (a synthetic, non-immunosuppressive derivative of FK506) increases EAAT2 protein levels and extends life span of ALS mice (Ganel et al. 2006; Rothstein et al. 2005). In addition, EAAT2 is also expressed by other cell types than astrocytes alone (Anderson and Swanson 2000), which were not targeted with this genetic experimental design. The beneficial effect of EAAT2 is often used as an explanation of beneficial effects found by cell transfers in ALS model rodents. For instance, the systemic transplantation of c-kit-positive cells from bone marrow in mutant SOD1 mice increases survival, which is, at least in part, attributed to increased EAAT2 expression induced by the transferred cells (Corti et al. 2010). The same holds true for the prolonged survival of ALS rats when treated with focal transplantation-based astrocyte replacement with wild-type glial-restricted precursors (GRPs) (Lepore et al. 2008). This study, interestingly, also focussed on the precise role of EAAT2 by also transplanting EAAT2 overexpressing GRPs and EAAT2 null GRPs. ALS mice treated with EAAT2 overexpressing GRPs show no additional increase of life span compared to wild-type GRP-treated ALS mice (which is already increased compared to controls). Intriguingly, this positive effect of transplantation of the wild-type GRPs is diminished in the mice transplanted with the EAAT2 null GRPs (Lepore et al. 2008). In addition, cocultures of human adipose-derived stem cells with astrocytes induce higher levels of EAAT2 in the astrocytes (Gu et al. 2010), though this treatment has not (yet) been shown to effect motor neuron survival in vitro or in vivo.

#### 4 Conclusion

Excitotoxicity plays a major role in the pathophysiology of ALS. Excitotoxins can cause neurodegeneration and ALS/PDC. In addition, the only drug that prolongs survival in patients targets excitotoxicity. The specific vulnerability of motor neurons to excitotoxic insults may be explained by their poor calcium buffering capacity and their expression of calcium-permeable receptors. Cell death in ALS can be postponed by blocking the AMPA receptor, and applying AMPA receptor agonists induces neuronal death. The AMPA receptor subunit responsible for calcium impermeability, GluR2, is less expressed by motor neurons in comparison to other neuronal subtypes, and knockdown of this subunit prolongs survival in ALS mice. Moreover, the pre-RNA editing to obtain normal GluR2 is diminished in motor neurons and in ALS patients. In accordance, increasing unedited GluR2 in mice induces neurodegeneration. Not only AMPA receptor calcium permeability but also glutamate transport is affected in ALS. Decreased levels of EAAT2 are present in ALS patients and ALS rodents, implying an increased vulnerability to glutamate. In addition, increased levels of EAAT2 prolong survival, and decreased levels reduce survival in ALS mice. Future research may focus further to develop therapeutic strategies targeting excitotoxicity in ALS.

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# **Excitotoxicity and Axon Degeneration**

# Anna E. King and James C. Vickers

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#### Abstract

Excitotoxicity has been implicated as a key pathogenic pathway in a number of neurodegenerative diseases and conditions including Alzheimer's disease, amyotrophic lateral sclerosis, multiple sclerosis, brain trauma, and stroke. While acute excitotoxicity can result in the initiation of cell death pathways, chronic or low levels of excitotoxin exposure may result in a more slowly progressing pathological cascade. In this respect, there is emerging evidence that excitotoxicity can result in axonal degeneration and pathology, a key pathological feature of many of these neurodegenerative conditions. Recent evidence supports the notion that axon degeneration can be a separate and independent

A.E. King  $(\boxtimes) \bullet$  J.C. Vickers

Wicking Dementia Research and Education Centre, University of Tasmania, Hobart, Tasmania, Australia

e-mail: kingae@utas.edu.au; James.Vickers@utas.edu.au

process from cell death, and thus mechanisms involved need to be understood in order to provide axonal protection in neurological disease. While axon degeneration following transection (Wallerian degeneration) has been well documented, less is known about axon degeneration following other insults such as excitotoxicity and the mechanistic relationships they bear to Wallerian degeneration. In particular, how a primarily somatodendritic insult, such as excitotoxicity, results in a pathological cascade within the axon is unclear. This chapter reviews our current understanding of the pathological changes and mechanisms of excitotoxin-induced axon degeneration and potential mechanisms involved. An increased understanding of the mechanisms of axon degeneration in neurological disease is essential to the development of therapeutic agents targeting axon protection.

List of Abbreviations					
Aß	Beta amyloid				
ALS	Amyotrophic lateral sclerosis				
AMPA	Alpha-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid				
CaMKII	Ca <sup>2+</sup> /calmodulin-dependent protein kinase II				
Cdk5	Cyclin-dependent kinase 5				
CNP	2'3'-Cyclic nucleotide 3'-phosphodiesterase				
CNS	Central nervous system				
CRMP2	Collapsin response mediator protein 2				
MAG	Myelin-associated glycoprotein				
MPP	1-Methyl-4-phenylpyridine				
NAD	Nicotinamide adenine dinucleotide				
NFH	Neurofilament heavy chain				
NFL	Neurofilament light chain				
NFM	Neurofilament medium chain				
NMDA	N-methyl-D-aspartate				
Plp	Proteolipid protein				
PNS	Peripheral nervous system				
TTX	Tetrodotoxin				
Wlds	Wallerian degeneration slow				

#### 1 Introduction

The toxicity of excitatory amino acids such as glutamate to neurons ("excitotoxicity") has been known since the 1950s (Lucas and Newhouse. 1957) and has since been implicated in the pathogenesis of a number of neurodegenerative conditions (Faden et al. 1989; Aarts et al. 2002; Hynd et al. 2004; Bogaert et al. 2010). Excitotoxicity was demonstrated by Olney (1969) to occur through overactivation of excitatory amino acid receptors and, in a variety of disease states, may also occur through the

impaired ability of cells to deal with sub-toxic levels of glutamate. In this regard, a characteristic of most neurodegenerative diseases is the slowly progressing cellular pathology and degeneration that occurs prior to overt cell death. Although excitotoxicity is generally considered a somatodendritic insult, there is also emerging evidence that excitotoxicity can play a role in axon degeneration, the latter a key feature of a number of neurodegenerative diseases including Alzheimer's disease, multiple sclerosis, and amyotrophic lateral sclerosis (ALS). The evidence for excitotoxic "axonopathy" and proposed mechanisms are discussed in this chapter.

# 2 Axon Degeneration in Neurodegenerative Disease

Axon degeneration is a prominent feature of neurodegenerative disease (Vickers et al. 2009) pathologically characterized by axonal swelling, axonal dystrophy, and distal axon dieback. It potentially represents one of the earliest findings in many diseases, and it has been proposed that it may be the cause of cellular dysfunction and degeneration through a dying back process. ALS is characterized by substantial early axonal degeneration in both the upper and lower motor neurons pathologically presenting as degeneration of the corticospinal tract and disconnection of the neuromuscular junctions and large axonal swelling referred to as "spheroids" in the proximal axons of lower motor neurons (Wood et al. 2003; Fischer et al. 2004). Causes of axonal degeneration have not been fully determined but are thought to involve axonal transport disruption. Axon transport disruption has also been implicated in axonal pathology in Alzheimer's disease, the latter which consists primarily of dystrophic neurites linked with Aß plaque formation (Vickers et al. 2009). Axon pathology linked with dystrophic neurite formation is accompanied by substantial demyelination (Mitew et al. 2010). Interestingly, putative demyelinating conditions, such as multiple sclerosis, are also associated with axonopathy and neuronal cytoskeletal changes indicative of axon transport disruption (Petzold et al. 2008). The precise cause of axonal degeneration in all of these conditions has not been determined and may be multifactorial in nature. However, there is substantial evidence for a role of excitotoxicity in the pathogenesis of these diseases, leaving the possibility that axonal degeneration results from excitotoxic processes.

#### 3 Mechanisms of Axon Degeneration

Axon degeneration following axonal transection has been very well defined and involves a process termed "Wallerian degeneration". A number of reviews describe the time course of Wallerian degeneration including the distinct phases of retraction of cut ends, formation of swollen end bulbs, latency period, axonal beading, sudden axonal fragmentation, and clearing of axonal segments by glial cells (e.g., Coleman 2005; Luo and O'Leary 2005; Wang et al. 2012,). Mechanistically, Wallerian degeneration has been shown to involve calcium influx (Knoferle et al. 2010), activation and loss of calpains (Glass et al. 2002; AraujoCouto et al. 2004),

activation of autophagy (Knoferle et al. 2010) and microtubule disruption (Zhai et al. 2003). Caspase activity and other apoptotic mechanism are not generally associated with Wallerian degeneration (Finn et al. 2000). Much of our understanding of the nature of axonal degeneration comes from a mutant mouse model, the slow Wallerian degeneration (Wlds) mouse, in which Wallerian degeneration is substantially delayed, suggesting that axon degeneration is an active process. Analysis of the genetic mutation that is responsible for this delayed degeneration revealed a splicing of two mRNAs to form a fusion protein consisting of Nmnat1 (an NAD+ salvage enzyme) and the N terminal protein of Ube4b (an E4 ubiquitin ligase) (Mack et al. 2001). Current research supports the notion that Nmnat1 activity is important for axonal protection in Wlds mice, and Ube4b may be responsible for directing it to the axon instead of the nucleus (Conforti et al. 2009). However, the protective role of Nmnat1 remains unclear (Sasaki et al. 2009, reviewed in Coleman and Freeman 2010). Despite the increasing knowledge on mechanisms of Wallerian degeneration, it remains unclear if other mechanisms of axonal degeneration exist. One type of axon degeneration that has been described that does not appear to involve Wallerian degeneration is developmental axon retraction, which occurs when a small shortening of the axon is required, for example, during neurite outgrowth and synapse formation. Axon retraction involves preservation of the axonal cytoplasm through gradual retrograde transport of materials to more proximal axon segments (reviewed in Luo and O'Leary 2005).

Axon degeneration that occurs in neurodegenerative disease, however, is not usually associated with direct severing of the axon, although it frequently shares many similarities with Wallerian degeneration, such as axon thinning, beading, and fragmentation (for review, see Saxena and Caroni 2007). In contrast, in a number of diseases and conditions, axon degeneration may occur by a dying back axonopathy, beginning at the distal axon (Cavanagh 1964), or may involve formation of large axonal swellings or spheroids (Trapp et al. 1998; King et al. 2011). Furthermore, the protective effect of the Wlds gene in some but not all (for review, see Coleman 2005) mouse models of disease suggests that different mechanisms may be involved. Pertinent to the current topic is whether axon degeneration induced by excitotoxicity occurs via a Wallerian-type degeneration or by a different mechanism.

# 4 Morphological Characteristics of Excitotoxin-Induced Axon Degeneration

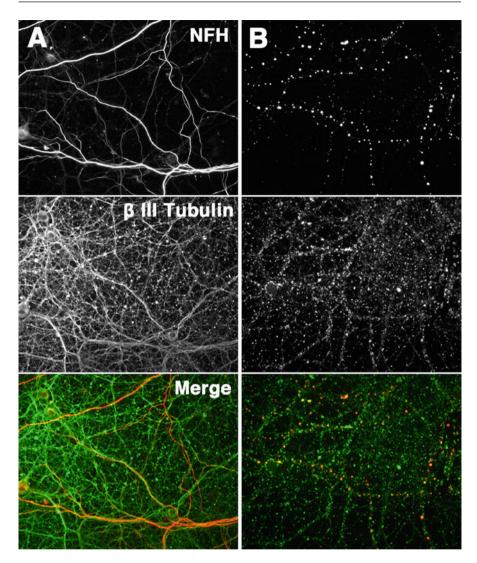
Despite its clear relevance to neurological diseases, few studies have examined the consequence of excitotoxicity to the axonal compartment. In vitro studies suggest that axonal changes may occur rapidly, prior to the hallmarks of neuronal cell death, and within 2–6 h of exposure to glutamate receptor agonists (King et al. 2007; Hou et al. 2009). Axonal degeneration also occurs following exposure to application of both NMDA (Chung et al. 2005; King et al. 2007) and non- NMDA (King et al. 2007) agonists. However, the type of degeneration occurring following such insults may be specific to both the particular pattern of receptor activation and the cell

type (King et al. 2007). In cultured rodent cortical neurons, glutamate, NMDA, and kainic acid exposure result in axonal retraction, beading, thinning, and subsequent fragmentation of the axon (Chung et al. 2005; King et al. 2007; Hou et al. 2009; Hosie et al. 2012) that resembles the morphological changes to axons undergoing Wallerian degeneration. However, cultured motor neurons may respond in an entirely different manner, particularly following kainic acid exposure, with a proportion of motor neuron axons displaying distinct axonopathic features characterized by large swollen structures in distal axon segments (King et al. 2007) that may show more resemblance to distal axon dieback described in a number of neurological diseases. These types of distal motor axon swellings have been described in mouse models of spinal muscular atrophy (Cifuentes-Diaz et al. 2002) although they may not be characteristic of all types of distal axon dieback, which in some cases may be more associated with distal axon atrophy and thinning (Schaefer et al. 2005). Distal axon degeneration was also more prominent than proximal degeneration in an in vivo rat model of somatodendritic NMDA exposure to retinal ganglion cells in a glaucoma model (Saggu et al. 2008, 2010).

# 5 The Role of the Cytoskeleton in Excitotoxin-Induced Axon Degeneration

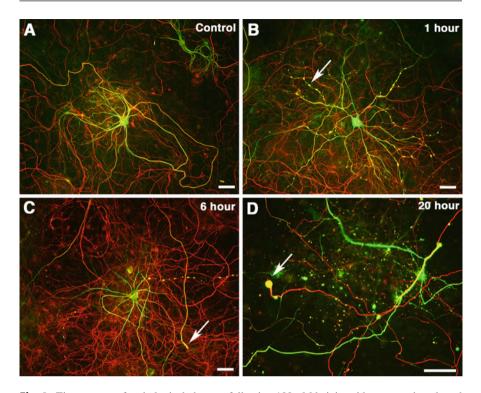
Axonal degeneration involves a breakdown of the components of the neuronal cytoskeleton. The neuronal cytoskeleton is composed of three distinct types of filaments, namely, microtubules, intermediate filaments (including neurofilaments), and microfilaments, all of which can undergo posttranslational modifications that affect their stability and function. Although distinct, the components of the cyto-skeleton are dynamically connected with each other, and there are a vast number of associated proteins for linking and regulation and to enable intracellular transport. Neurofilaments, consisting of light (NFL, 61 KDa), medium (NFM 90 KDa), and heavy (NFH 115 KDa) chains (Lariviere and Julien 2004), are fibrous subunits that are only present in some eukaryotic organisms including vertebrates, nematodes, and mollusks and are likely to have broader functions than just serve as a structural scaffold (reviewed in PerroneCapano et al. 2001).

Cytoskeletal breakdown has been previously characterized for classic forms of Wallerian degeneration. In the initial stages, there is a depolymerization of the *microtubules*, which occurs within 4 h in in vitro models (Zhai et al. 2003). Depolymerization of microtubules is not accompanied by tubulin degradation, but is followed at 8–12 h by loss of neurofilament subunits (Zhai et al. 2003). Both microtubule depolymerization and loss of neurofilaments are dependent on degradation by the ubiquitin-proteasome system (Zhai et al. 2003). Our studies have examined the cytoskeletal changes in cultured cortical neurons following excitotoxic stimulation following a range of glutamate concentrations. A chronic low-dose treatment resulted in specific loss of NFH, with a relative sparing of the NFM and NFL subunits at 24 h posttreatment (Fig. 1. Chung et al. 2005). While beading of NFM and NFL were apparent at this time point with higher glutamate



**Fig. 1** Cytoskeletal architecture in primary cultured rat cortical neurons at 21 DIV. (a) In untreated cultures NF-H immunoreactivity was found primarily in thick axonal processes, and co-localized with  $\beta$ III-tubulin. (b) Following treatment with 10  $\mu$ M glutamate fragmentation of microtubule proteins was present accompanied by beading and loss of neurofilament proteins, particularly the NFH subunit (Modified with permission from Chung et al. 2005)

treatments (100  $\mu$ m), their degradation was delayed relative to the NFH subunit. The differential vulnerability of neurofilament subunits may reflect their temporal expression during development, with the NFH subunit being the last to be expressed. In a similar fashion to Wallerian degeneration, tubulin immunoreactivity remained intact until later stages indicating that tubulin subunits were not degraded.



**Fig. 2** Time-course of pathological changes following 100  $\mu$ M kainic acid exposure in cultured motor neurons at 21 DIV. (a) Untreated motor neuron immunolabelled for dephosphorylated neurofilament protein (*green*) and phosphorylation-independent NFM (*red*). (b) Following 1 h of kainic acid exposure, beading was present in dendrites (*arrow*). (c) Axon swellings (e.g., *arrow*) developed following 6 h of kainic acid exposure and were frequently associated with expression of de-phosphorylated neurofilament protein (*green*) in the axonal network. (d) Following 20 h kainic acid exposure the distal portions of axonal segments frequently expanded to form large bulbous structures (*arrows*). Scale Bars: 50  $\mu$ m (Modified with permission from King et al. 2007)

Distal axon degeneration induced by low-dose kainic acid in cultured spinal motor neurons, on the other hand, was characterized by a distinct set of cytoskeletal changes (King et al. 2007). In these neurons, dephosphorylated neurofilament protein (SMI32 immunoreactivity) is normally localized to the somatodendritic compartment and the proximal axon (King et al. 2007). However, following kainic acid treatment, there was a mislocalization of non-phosphorylated neurofilament to the more distal axon compartment, which was accompanied by accumulation of neurofilament proteins and mitochondria in axonal swellings (Fig. 2). Swollen, neurofilament-rich distal axon segments may be caused by a disruption of retrograde axonal transport mechanisms and have been shown to occur in a mouse model of Kennedy's disease due to decreased expression of the retrograde transport motor protein dynactin 1 (Katsuno et al. 2006). The presence of multiple cytoskel-etal components and organelles in these swellings suggests a nonspecific disruption

of retrograde transport. Alterations in neurofilament phosphorylation have been reported in other models of kainic acid toxicity, with decreased phosphorylated and increased non-phosphorylated epitopes of NFH in a rat model of kainic acid exposure to the hippocampus (Wang et al. 1992) and in cultured spinal neurons exposed to sublethal kainic acid (Vartiainen et al. 1999). In contrast, blockade of non-NMDA receptors in cultured spinal neurons (Vartiainen et al. 1999) or glutamate exposure of dorsal root ganglion neurons (Kesavapany et al. 2007) caused increased neurofilament phosphorylation, which was localized to the soma. The significance of the changes in localization of these neurofilament epitopes remains unclear; however, dephosphorylation has been implicated in slowing of axonal transport through irregular and fused crossbridges (Gotow 2000). Conversely, in cultured cortical neurons, glutamate toxicity has been linked to slowing of anterograde transport through increased neurofilament phosphorylation and resultant dissociation of neurofilament from kinesin anterograde motors (Yabe et al. 2001). Thus, it is possible that both phosphorylation and dephosphorylation of neurofilament proteins can affect transport within the axon.

# 6 Molecular Mechanisms of Cytoskeletal Disruption in Excitotoxin-Induced Axon Degeneration (Calpains Versus Caspases)

The role of specific proteases in cytoskeletal breakdown following excitotoxic insult remains unclear. Calpains, calcium-activated cysteine proteases, have been implicated in cytoskeletal breakdown following traumatic brain injury and axon transection (Kampfl et al. 1996) and in focal axonal beading (Kilinc et al. 2009). Furthermore, a study by Higuchi et al. (2005) demonstrated the protective effect of calpain, but not caspase, inhibition on the degeneration of neuronal processes and downstream subacute cell death, following kainic acid exposure in vivo, although acute cell death was not affected. In this model, calpain activation was associated with increased tau phosphorylation through cleavage of the CDK5 activator p35 to p25 (Nath et al. 2000; Kilinc et al. 2009). Calpain activation was higher in the somatodendritic compartment leading to the hypothesis that tau phosphorylation occurs in the soma, resulting in depletion of axonal tau, with microtubule disruption-induced axonal degeneration occurring later than somatodendritic changes. Detailed analysis of the axonal processes in this model was not performed, and the role of calpain in axonal beading, a common feature of both excitotoxic and Wallerian axon degeneration, was not determined. However, in a cell culture model of glutamate excitotoxicity in cortical neurons, calpain inhibition was protective against axonal beading and thinning that occurred from 2 h post-glutamate exposure (Hou et al. 2009). The authors showed that, in addition to calpain, CaMKII is activated, and the two proteins affect the microtubule-associated protein (Fukata et al. 2002) collapsin response mediator protein 2 (CRMP2). CaMKII causes phosphorylation of CRMP2, which is protective against calpain-mediated proteolysis, although CaMKII inhibition is also protective against axonal beading, suggesting a central modulatory role of CRMP2 in axon integrity (Hou et al. 2009). Furthermore, overexpression of CRMP2 protects against axonal beading and subsequent degeneration. It is interesting to note that phosphorylation of neurofilaments has also been shown to protect these proteins from calpain-mediated proteolysis (Goldstein et al. 1987). Thus, dephosphorylation of neurofilaments, as occurs in motor neuron axons following kainic acid exposure (King et al. 2007), may make these proteins more vulnerable to proteolysis. Neurofilament proteolysis is involved in removing excess neurofilaments under physiological conditions (reviewed in Liu et al. 2004), and thus, it is unclear whether dephosphorylation of neurofilament is a protective mechanism to remove excess neurofilament from the axon terminal following transport disruption or a consequence of the pathological cascade (King et al. 2007).

Caspase activation, which can occur by a number of different mechanisms including both intrinsic pathways such as mitochondrial release of cytochrome c and extrinsic pathways such as activation of death receptors, is thought not to be involved in axon degeneration following transection (Wallerian degeneration) despite its activation in the soma (Finn et al. 2000). Similarly, in models of neurodegenerative disease such as ALS, caspase inhibition fails to protect against axon degeneration despite protection from neuronal cell death (Gould et al. 2006). Conversely, caspase activation has been implicated in axon degeneration following a number of insults including psychosine toxicity (a sphingolipid involved in Krabbe's disease, Smith et al. 2011), amyloid toxicity (Nikolaev et al. 2009), and MPP toxicity (Morfini et al. 2007). Reports of the role of caspase activation following growth factor deprivation are controversial, with some authors finding that it is involved (Schoenmann et al. 2010) and others finding no involvement (Finn et al. 2000). Our studies using cortical neurons in compartmented chambers to allow application of toxins and inhibitors to either the axon or somatodendritic compartment of the cell indicated that caspase inactivation with the pan-caspase inhibitor Z-FAD-FMK to either the somatodendritic compartment or the axon provides some protection against axon degeneration induced by somatodendritic excitotoxicity, with an overall decrease in axon degeneration of 42 % and 33 %, respectively (Hosie et al. 2012). This suggests that axon degeneration following excitotoxicity involves degenerative processes occurring in both the axon and the soma. Elucidating the role of calpains and caspases in excitotoxic axon degeneration in specific cell types is an area of future investigation.

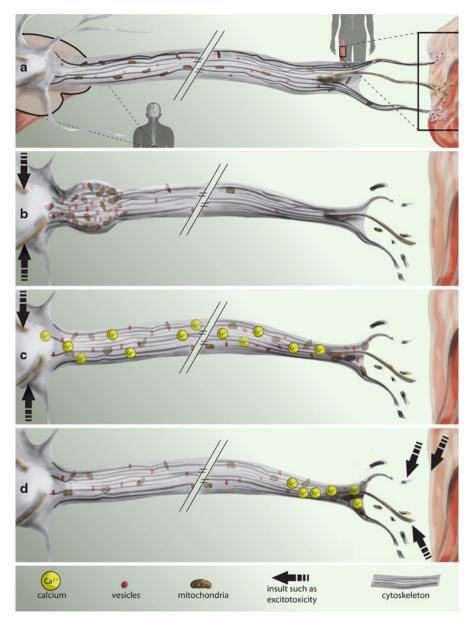
# 7 How Is Degeneration Initiated in the Axon Following Excitotoxic Insult?

Following excitotoxic insult, degeneration is initiated in the axon, and the type of degeneration is cell type specific. However, it remains unclear how the insult is transferred from glutamate receptors, the mediators of the insult, to the axonal

compartment. Glutamate receptors involved in synaptic signaling are predominantly located on the postsynaptic densities of glutamatergic synapses and at extrasynaptic sites on the somatodendritic compartment of neurons. Increase in calcium influx in this region is unlikely to spread to the axon terminals. It is possible that axon degeneration results only from cell death; however, as discussed, protection from cell death does not protect the axon, and axon degeneration is apparent prior to cell death (Chung et al. 2005; Higuchi et al. 2005; King et al. 2007; Hou et al. 2009). Axon degeneration may be initiated through a number of different mechanisms that will be examined in more detail: (1) nonapoptotic somatic changes resulting in axon transport disruption and subsequent axon starvation, (2) soma-axon signaling, and (3) glutamate receptor expression on the axonal compartment (Fig. 3).

#### 7.1 Axon Starvation

A potential mechanism of initiation of axon degeneration through excitotoxic insult is through transport disruption, which results in reduced delivery of mitochondria or cytoplasmic components to the axon. Slowing of the transport of neurofilament proteins has been demonstrated following glutamate exposure to primary cortical neurons in culture (Ackerley et al. 2000). Transport disruption was accompanied by an increased phosphorylation of the neurofilament subunits (Ackerley et al. 2000), which has previously been associated with decreased transport rates (Lewis and Nixon 1988). Accumulation of phosphorylated neurofilament subunits in the soma of dorsal root ganglia following exposure to glutamate has also been reported (Kesavapany et al. 2007). As described earlier, aberrant phosphorylation of microtubule-associated proteins such as tau may prevent their transport into the axon leading to microtubule disassembly. The differential role of phosphorylation of cytoskeletal proteins in the soma in axon degeneration in different neuronal populations has not been investigated. More recently experiments in cultured hippocampal neurons have demonstrated disruption of motor proteins themselves following excitotoxic stimulation. A C-terminally truncated form of the retrograde dynein/dynactin motor complex p150Glued was formed after glutamate exposure and was also present in tissue from Alzheimer's disease cases (Fujiwara and Morimoto 2012). Overexpression of the wild-type protein was protective against neuritic beading and knockout of the protein exacerbated beading (Fujiwara and Morimoto 2012). Collectively, these data suggest a role for transport failure in excitotoxin-induced degeneration; however, the mechanisms by which axon transport failure results in axon degeneration remain unclear. Recent data indicates that axon transport disruption alone may not be sufficient to induce axon degeneration, at least on a rapid scale (Marinkovic et al. 2012). Furthermore, the links between failure of axon transport in the soma and axonal blebbing or distal axonal neurofilament accumulations, which occur rapidly following excitotoxin exposure in cell culture models, remain to be elucidated.



**Fig. 3** Potential mechanisms of axon degeneration induced by excitotoxicity. (a) Motor axons are spatially separated from the somatodendritic compartment. (b) Excitotoxicity to the somatodendritic compartment may result in transport blockage and withering of the distal axon. (c) Somatodendritic exposure may result in a signal (such as calcium, or newly translated protein) being transferred to the distal axon regions. (d). Distal axon exposure to excitotoxicity may directly result in degeneration (Figure courtesy of Graeme McCormack)

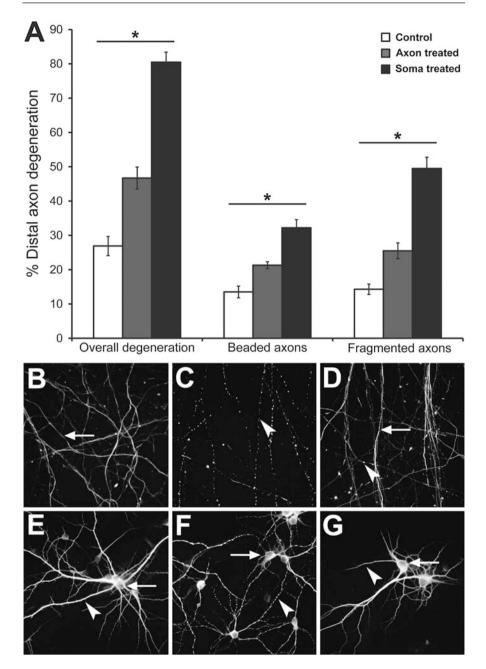
#### 7.2 Axon-Soma Signaling

An alternative mechanism of initiation of axon degeneration following somatodendritic glutamate receptor stimulation is production of an axon destruction signal and/or loss of an axon maintenance signal. There is little information on this type of initiation of axon destruction instigated specifically by excitotoxicity. However, excitotoxic mechanisms of degeneration are calcium dependent; therefore, studies examining changes in calcium influx following nontoxic glutamate exposure may give an indication of changes that could occur following excitotoxicity. A study by Yamada and colleagues (2008), in developing granule neuron cultures, demonstrated that growth cone modulation by glutamate in immature neurons occurs through perikaryal AMPA receptormediated glutamate exposure, suggesting soma to distal axon signaling. Despite the absence of action potentials in these immature neurons, signaling was the result of a sweep of calcium along the axon. Both perikaryal and distal axon glutamate exposure resulted in a fast calcium transient throughout the whole cell, but only perikaryal exposure resulted in a subsequent calcium sweep from proximal to distal axon. The calcium sweep could also be initiated by depolarization and perikarval calcium increase and involved release of calcium from intracellular stores. A study by Christie and Jahr (2008) examined the contribution of NMDA receptor stimulation to axonal calcium increase in stellate cells. They demonstrated that bath application of NMDA to cerebellar slice cultures, in the presence of TTX to block action potentials, caused increases in intracellular calcium concentrations in both axons and dendrites. Axonal calcium increases particularly occurred at axonal boutons, thought to be the sites of en passant presynaptic terminals. However, calcium increase was higher in boutons closer to the soma and lower in distal axon segments, suggesting that the increase in calcium arose from somatic NMDA receptor activation, with a passive propagation of depolarization to axonal voltage-gated calcium channels. Local activation of dendritic NMDA receptors confirmed this somatodendritic transfer (Christie and Jahr 2008). In addition to passive transfer of calcium currents, action potentials caused large increases in axonal calcium concentrations in boutons (Christie and Jahr 2008). It is currently unclear if these calcium increases are sufficient to cause activation of axonal degeneration mechanisms or if axonal beading corresponds to positions of boutons in axons. However, together these data suggest that somatodendritic glutamate exposure can result in axonal calcium influx even without action potential. In the event that axon potentials are initiated in hyperexcitable neurons, it has been shown that calcium entry can occur through the reverse operation of the sodium calcium exchanger that can be the result of persistent sodium influx (Stys et al. 1992).

Aside from calcium influx, studies investigating other intracellular signaling mechanisms in excitotoxin-induced axon degeneration are lacking. In terms of axon degeneration, inhibition of protein synthesis in the soma results in axon degeneration without cell loss and proteasome inhibition following axon transection is protective against axon degeneration, supporting the hypothesis that a protein/proteins synthesized in the soma are required for axon maintenance (Gilley and Coleman 2010). These authors have identified the NAD+ synthesizing enzyme Nmnat2 (an isoform of the Wlds protein Nmnat1) as a likely candidate for an axonal survival signal. Involvement of these processes in excitotoxin-induced axon degeneration and distal axon dieback has not yet been investigated.

#### 7.3 Axonal Expression of Glutamate Receptor Proteins

Although the prime site of glutamate receptor expression is on the postsynaptic density of the somatodendritic compartment, where they are involved in synaptic transmission, recent advances have demonstrated other roles for glutamate receptors (Araque and Perea 2004), and their expression has been demonstrated at a number of non-synaptic locations. Their presence at extrasynaptic sites on the somatodendritic compartment has been well documented (Passafaro et al. 2001; Tovar and Westbrook 2002; Kane-Jackson and Smith 2003) where they are thought to play a modulatory role (Diamond and Jahr 2000; Jourdain et al. 2007) in addition to acting as a reserve supply for synaptic receptors (for recent review, see Newpher and Ehlers 2008). The axonal expression of glutamate receptors has been less well documented, although they have been reported to be present in immature neurons (Yamada et al. 2008) on growth cones (Zheng, et al. 1996; King et al. 2006) and are present in presynaptic terminals (recently reviewed in Pinheiro and Mulle 2008). Recently AMPA/kainate receptors have been demonstrated to be present on myelinated axons (Ouardouz et al. 2009a, b). The functional activity of axonal receptors is currently disputed (e.g., Berretta and Jones 1996; Huang and Bordey 2004; Christie and Jahr 2008). In many excitotoxic models, particularly in cell culture, excitotoxins are bath applied, leaving the possibility that excitotoxin-induced axon degeneration occurs directly through stimulation of axonal glutamate receptors. To directly test the hypothesis that axonal glutamate exposure can result in axon degeneration, we have recently utilized compartmented microfluidic chambers (Taylor et al. 2005) to allow specific exposure of the axonal compartment to glutamate. Our findings demonstrate that axon degeneration characterized by beading and fragmentation resulted from prolonged (24 h), but not acute (Underhill and Goldberg 2007), exposure of the axons to glutamate without subsequent apoptosis of the soma, even after 72 h (Fig. 4, Hosie et al. 2012). Western blot analysis confirmed the presence of AMPA/kainate receptors in the axons, although NMDA receptors could not be detected by this method. In this study, caspase inhibition was protective against axonal excitotoxin-induced axon degeneration. Interestingly protection was higher when caspase activation in the soma was prevented, suggesting that axon-soma-axon signaling is involved in degeneration rather than direct degeneration resulting from calcium influx in the axon (Hosie et al. 2012).



**Fig. 4** Axon degeneration in cortical neurons grown in compartmentalized microfluidic chambers (Taylor et al. 2005) and exposed to 100  $\mu$ M glutamate to either the somatodendritic compartment or the isolated axon compartment. (a) Quantitation of axonal beading and fragmentation following glutamate exposure indicated a significant threefold increase in axon degeneration (beading plus fragmentation) following somatodendritic exposure, and a significant 1.5-fold increase following

#### 8 White Matter Excitotoxicity

In addition to excitotoxicity directly acting on the neuron to induce axon degeneration, there is increasing evidence that axon degeneration can be initiated in white matter indirectly through disturbed neuron-glia interactions. White matter tracts are vulnerable to damage following brain trauma (Kraus et al. 2007; Sharp and Ham 2011) and hypoxia/ischemia (Pantoni et al. 1996; Yam et al. 1998), and white matter excitotoxicity has been particularly implicated in multiple sclerosis (Pitt et al. 2000; Smith et al. 2000). Direct application of excitotoxins to white matter tracts of the optic nerve (Matute 1998) or internal capsule (Fowler et al. 2003) confirmed the role of excitotoxicity in the degeneration of the white matter tracts as a distinct process from somatodendritic exposure. Damage is characterized by injury to both the axonal process and surrounding glial cells. Axonal damage included swollen and bulbous profiles, cytoskeletal disruption such as loss of neurofilament (Fowler et al. 2003; Cuthill et al. 2006), and disruption to axonal electrical activity (Stys et al. 1992). Nonneuronal damage included damage to the myelin sheath (Li and Stys 2000) and focal swelling and cytoskeletal breakdown of oligodendrocytes and astrocytes (Pantoni et al. 1996; Li and Stys 2000). Although damage to axons following hypoxia can occur through non-excitotoxic pathways (Underhill and Goldberg 2007), non-NMDA (Agrawal and Fehlings 1997; Li and Stys 2000; Tekkok and Goldberg 2001) receptor antagonists are protective against white matter degeneration both in vitro and in vivo, suggesting a role for excitotoxicity in this damage, specifically mediated through non-NMDA receptors. The role of NMDA receptors remains less clear (Yam et al. 2000; Li and Stys 2000), although recent reports similarly suggest a role for NMDA receptor stimulation (Bakiri et al. 2008). In acute spinal cord slices, white matter degeneration occurs via a calcium-dependent mechanism (Li and Stys 2000).

White matter excitotoxicity is thought to occur predominantly through *oligodendrocytes*, which possess both AMPA and NMDA receptors, localized to the soma and the myelin sheath (Li and Stys 2000; Brand-Schieber and Werner 2003; Karadottir et al. 2005; Salter and Fern 2005; Micu et al. 2006) and have been shown to be vulnerable to excitotoxicity both in vitro (Yoshioka et al. 1996; Sanchez-Gomez and Matute 1999) and in vivo (Matute 1998). Toxicity can occur both through direct stimulation of the receptors and indirect mechanisms involving transporters (Oka et al. 1993), and the mechanisms involved are dependent on the type and duration of insult. For a recent review on white matter excitotoxicity and

**Fig. 4** (continued) exposure to the isolated axon compartment. Representative immunocytochemical images of axons (NFM, **b**–**d**) and somatodendritic compartment (MAP2, **e**–**g**) in control (**b**, **e**), soma treated (**c**, **f**) and axon treated (**d**, **g**) cultures. Axons demonstrated increased beading and fragmentation following treatments. *Arrows* indicate whole axons, arrowheads indicate fragmented and beaded axons. MAP2 immunoreactive dendrites demonstrated increased beading following somatodendric exposure (**f**) but not following axonal exposure (**g**). *Arrows* indicate soma, arrowheads indicate dendrites. Scale bar: 50 µm (Reproduced with permission, from Hosie et al. 2012)

its role in disease, see reviews by Matute et al. 2007; Matute and Ransom 2012. The question remains to as to how oligodendrocyte damage following glutamate exposure results in axonopathy.

Recent advances have detailed interactions between oligodendrocytes and axons that extend far beyond the role of passive insulation. Signaling between the myelin sheath and the axon is essential in maintaining fine-tuning of myelination, which is important in regulation of precise neuronal activity within a network (Wake et al. 2011). A type of axo-myelinic synapse has been proposed with glutamate receptors on both the axon and innermost compact myelin sheath surrounding the axon (Stys 2011). It is hypothesized that this type of myelin-axon signaling could be responsible for activity-dependent signaling that allows optimization of axonal sheath thickness. In terms of axon support, it has been proposed that oligodendrocytes and other glia (Magistretti 2006) may provide neurotrophic support to the axon in times of energy shortage from the soma (reviewed in Nave 2010). Specific myelin/oligodendrocyte proteins additionally may promote axonal integrity and survival. Mouse models with knockout of the oligodendrocyte proteins plp and CNP have frequent axonopathy characterized by axonal swelling and cytoskeletal disruption (Lappe-Siefke et al. 2003; Edgar et al. 2004). These axon disruptions occur in the absence of gross myelin loss highlighting the importance of specific proteins in myelin-axon signaling to axonal maintenance and function. Myelin has been shown to have a protective effect on axons in injury paradigms and is protective against mild stretch injury (Staal and Vickers 2011). This protection may be through myelin-associated glycoprotein (MAG) signaling to the axon, as knockout of MAG in mice results in increased axon degeneration in both the CNS and PNS and increased axonal vulnerability to insults such as acrylamide toxicity (Nguyen et al. 2009). Protective effects of MAG may be through cytoskeletal stabilization as it has been shown to affect the phosphorylation of neurofilament heavy and medium chains (NFH and NFM), which in turn affects the caliber of the axon (Hsieh et al. 1994; Dashiell et al. 2002; Garcia et al. 2003). Myelination and specific myelin protein may also affect axon transport mechanisms. Plp has been shown to play a role in axonal transport (Edgar et al. 2004).

The importance of demyelination and loss of glial support of the axon is highlighted in multiple sclerosis, in which immune attack of the myelin sheath results in axonopathy (reviewed in Dutta and Trapp 2011). However, the role of excitotoxic mechanisms in white matter injury remains an area for future research. Axon degeneration may result directly from loss of interaction with myelin, specific oligodendrocyte proteins, loss of neurotrophic support, or alternatively direct excitotoxicity on the axon shaft (Hosie et al. 2012).

#### 9 Conclusion

A number of diseases are characterized by axon degeneration, and excitotoxicity has been implicated as a pathogenic mechanism in these diseases. Axon degeneration may occur through a number of different excitotoxic mechanisms including somatodendritic glutamate exposure, direct axonal glutamate exposure, and glial glutamate toxicity. A better knowledge of the mechanisms involved in specific diseases may aid in directing therapeutic interventions. For example, excitotoxicity is implicated in both multiple sclerosis and ALS, and both have evidence of increased glutamate within the cerebrospinal fluid (Shaw et al. 1995; Barkhatova et al. 1998) and involve axon degeneration. However, the mechanisms of axon degeneration are likely to be different, with ALS potentially involving somatodendritic exposure to the spinal motor neurons and subsequent axonal degeneration and multiple sclerosis involving degeneration through loss of myelin proteins. Therapeutics may need to target specific complements of glutamate receptors, or downstream signaling pathways, that are expressed on the axon, on glial cells, or at synaptic or non-synaptic sites on neurons.

An alternative mechanism of therapeutic intervention would be to target the pathways of axonal degeneration, which may show common features between various disorders. Cytoskeletal degeneration, in particular, is an attractive target, and trials of drugs that stabilize the microtubule network, such as taxol and epothilone, are having positive outcomes in a number of mouse models involving axonopathy including tau transgenic mouse models of AD pathology (Brunden et al. 2010; Barten et al. 2012; Zhang et al. 2012) and spinal cord injury (Hellal et al. 2011). In either scenario, drug targets that are well defined to specific components of the pathological cascade leading to axonal and neuronal degeneration are more likely to lead to successful treatment strategies. Thus, increasing our knowledge of mechanisms of axon degeneration in specific neurodegenerative conditions is fundamental to developing appropriate therapeutic agents.

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## **Excitotoxicity in HIV Associated Neurocognitive Disorders**

## Belinda Cruse and Bruce J. Brew

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#### Abstract

HAND (HIV associated neurocognitive disorders) represents a group of disorders, encompassing HIV associated asymptomatic neurocognitive impairment (ANI), HIV-associated mild neurocognitive disorders (MND), and HIV associated dementia (HAD). Even in the era of highly-active antiretroviral therapy, the prevalence of HAND is thought to be around 40 %, with milder forms making up a greater proportion of cases. HAND are clinically manifest as subcortical dementing

B. Cruse

B.J. Brew (🖂)

University of New South Wales, Sydney, Australia

Department of Neurology, St. Vincent's Hospital, Sydney, NSW, Australia e-mail: belindacruse@hotmail.com

St. Vincent's Centre for Applied Medical Research, Darlinghurst, NSW, Australia

Departments of Neurology and HIV Medicine, St. Vincent's Hospital, Sydney, NSW, Australia e-mail: B.Brew@UNSW.edu.au

processes; correlating with the preferential involvement of the basal ganglia, subcortical white matter, frontal lobe and hippocampus in neuropathological studies.

A central mechanism leading to synaptic dysfunction and loss, neuronal dysfunction and apoptosis in HAND is excitotoxicity mediated through the excessive activation of neuronal (and probably astrocytic) NMDA receptors (NMDAR). There is direct and indirect evidence that in HAND, increased neuronal NMDAR activation occurs due to elevated extracellular glutamate; a result of both increased presynaptic release and impaired clearance from the synaptic space. In addition, high levels of quinolinic acid (QUIN) (an NMDAR agonist) released by activated macrophages (and to a lesser extent microglia) lead to excessive NMDAR activation and excitotoxicity. Furthermore, HIV related proteins (particularly gp120 and Tat) can both activate the NMDAR, and augment receptor activation by glutamate and other agonists.

In vitro evidence supports the role of NMDAR antagonists in the prevention of neuronal loss induced by HIV related proteins. Clinical studies have demonstrated that the NMDAR antagonist memantine is safe and tolerable in HAND patients; so far clinical trials have failed to demonstrate a clinically beneficial effect of this therapy on neurocognitive function in HAND patients, however trials are ongoing.

#### **Keywords**

Dementia • Excitotoxicity • Glutamate • HIV • Neurocognitive disorders • NMDAR

### 1 Introduction

This chapter focuses the reader on the role of excitotoxicity in the pathogenesis of HIVassociated neurocognitive disorders (HAND) after first considering the epidemiology of this group of conditions and briefly outlining the pathological and clinical features of HAND. An overview of the multiple neuropathogenic pathways that contribute to HAND is presented, with a particular emphasis on changes in clinical, pathological, and possibly pathophysiological aspects of HAND that are taking place in the era of highly active antiretroviral therapy (HAART). Evidence for, and mechanisms of, NMDA receptor (NMDAR)-mediated excitotoxicity in HAND is then described in detail, followed by a discussion of current evidence regarding the use of adjunctive therapies (and more specifically NMDAR antagonists) in the treatment of HAND.

## 2 Epidemiology of HAND

HAND represents a group of disorders, encompassing HIV-associated asymptomatic neurocognitive impairment (ANI), HIV-associated mild neurocognitive disorders (MND), and HIV-associated dementia (HAD) (Antinori et al. 2007). Despite the considerable prognostic improvement achieved in the HIV-infected population with the introduction of highly active antiretroviral therapy (HAART), it so far has failed to fully protect treated individuals from HAND, even in those patients in whom virologic suppression is achieved in the plasma and CSF (Simioni et al. 2010; Robertson et al. 2007; Cysique and Brew 2011; Cysique et al. 2006). Epidemiological studies in the post-HAART era indicate the prevalence of these disorders at around 40 % (similar to that in the pre-HAART era), despite a decrease in the prevalence of HAD after HAART introduction (Cysique et al. 2004; Heaton et al. 2010; Dore et al. 1999). Therefore, while milder forms of HAND (ANI and MND) make up a greater proportion of cases, these disorders impact significantly on affected individuals' quality of life and independence (Antinori et al. 2007) and mortality (Ellis et al. 1997) and clearly impact socioeconomically. HAND also appears to impact HIV-infected individuals at higher CD4 T-cell counts in the post-HAART era (Sacktor et al. 2001; Brew 2004).

## 3 Pathological Features

HIV probably enters the central nervous system (CNS) via infected monocytes from the periphery. These cells cross the blood-brain barrier (BBB) both as part of the usual process of perivascular macrophage turnover and because the permeability of the BBB is pathologically increased in the context of HIV infection (by the effects of systemic inflammation and the direct effects of gp120 and tat proteins (reviewed by Ivey et al. 2009; Langford and Masliah 2001; Toborek et al. 2003; Annunziata 2003)). While both of these processes are shown to contribute to the entry of HIV into the CNS, it is currently unclear which factor is of greater importance in the early pathogenesis of HAND or in fact which occurs first. Furthermore, transmigration across the BBB by infected and uninfected monocytes and macrophages is accelerated by inflammatory cytokines (produced in the periphery and centrally, as a result of neuroinflammation) (Persidsky et al. 2000), with resultant amplification of monocyte/macrophage chemotaxis and HIV delivery to the CNS. Within the CNS HIV productively infects monocytes and microglial cells and infects a smaller proportion of astrocytes in a restricted, non-cytolytic state (Churchill et al. 2009; Wang et al. 2004; Thompson et al. 2004).

Pathological changes, neuronal dysfunction and apoptosis, and corresponding volume loss on neuroimaging preferentially affect subcortical deep grey structures, the hippocampus, and the frontal cortex (Lindl et al. 2010; Bell 2004; Boissé et al. 2008; Moore et al. 2006).

Microscopically, HIV infection is associated with inflammatory infiltrates of mononuclear cells, with microglial and astrocyte activation. The neuroinflammatory changes in HIV infection are termed encephalitis (HIVE). Infiltration of bloodderived macrophages, together with microglial cells and, virtually pathognomonic of HIVE, multinucleated giant cells, are seen. Microglial nodules, multinucleated giant cells, and HIV antigens are frequently located in the basal ganglia and in a subset of patients in the subcortical white matter (termed HIV leukoencephalitis) (Petito 2003). There is also a widespread reactive astrocytosis and a variable degree of myelin damage (Bell 2004) and a sparse lymphocytic infiltrate in leptomeninges and perivascular spaces (Petito 2003). Neurons are not infected by HIV; however, they are clearly affected, with diffuse or focal neuronal cell death in the grey matter in severe cases (Petito 2003). Furthermore diffuse synaptic and dendritic loss, together with neuronal loss, is reported in HIV-infected individuals (Everall et al. 1999; Bell 2004).

The clinical features that predominate in HAND are those of a subcortical dementing process, with effects on cognitive slowing and impairment, behavioral and emotional change (usually apathy, especially in the HAD stage), together with prominent gross and fine motor slowing and gait disturbance (McArthur et al. 2005; Staekenborg et al. 2008). While currently the exact clinicopathological correlate is not fully understood in HAND (Gray and Keohane 2003), neuronal degeneration (i.e., neuronal degradation and death) underlies the neuropsychological impairment in HIV infection (Moore et al. 2006). Massive neuronal death, however, is not commonly seen and is not the main pathological counterpart of HAND (Adle-Biassette et al. 1999). In fact milder forms of HAND appear to correlate more closely with reduced synaptic and dendritic density rather than overt neuronal loss (Everall et al. 1999). Importantly, some of these changes are thought to be reversible (Gray and Keohane 2003), offering potential therapeutic targets.

## 4 Neuropathogenesis of HAND

Neuronal and astrocyte loss in HAND have historically been found in close anatomical proximity to the aforementioned inflammatory infiltrates and activated microglia (Brew et al. 1995; Adle-Biassette et al. 1999; Kaul et al. 2005). These neuropathological findings imply that neurotoxicity in HAND occurs through indirect mechanisms, specifically through toxins produced by activated mononuclear cells and microglia, rather than as a direct effect of HIV. Further supporting this widely held view is the fact that cognitive changes observed clinically in HAND are more closely associated with neuronal loss, decreased synaptic and dendritic density, intensity of local inflammation (such as the number of microglial cells and markers of tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) expression by microglia and astrocytes), and evidence of excitotoxins, as opposed to the number of HIV-infected cells or concentration of HIV proteins (reviewed in Kaul et al. 2001; Adle-Biassette et al. 1999).

Monocytes and microglial cells are activated by HIV infection; exposure to extracellular HIV proteins, cytokines, and chemokines liberated from other similarly activated microglial cells (reviewed by Bol et al. 2011); and other factors including quinolinic acid (QUIN), glutamate and L-cysteine, arachidonic acid, platelet-activating factor (PAF), NTox, superoxide anions, and TNF- $\alpha$ . Monocytes and microglial cells are considered central to the pathogenesis of HAND, both by virtue of their role as the principle cells infected by HIV and as a major source of neurotoxins (Valle et al. 2004). Importantly, however, abnormalities of astrocyte

function due to neurotoxic actions of some of the above cytokines and chemokines, astrocyte activation and astrocytosis, and astrocyte loss have also been implicated (Kaul et al. 2001) and will be further discussed. Indeed, while there is evidence of a direct role of some HIV proteins in the neuropathogenesis, the observed dissociation between the HIV viral burden in the brain and clinical deficits points to relatively more important roles of toxins produced by activated monocytes and microglial cells, most of which are uninfected with HIV (Brew et al. 1995).

A central mechanism leading to synaptic dysfunction and loss, neuronal dysfunction, and apoptosis in HAND is excitotoxicity mediated through the excessive activation of neuronal (and probably astrocytic) NMDAR and the subsequent, stereotyped cascade of intracellular events driven by intracellular Ca<sup>2+</sup> overload and shared by many neuroinflammatory and neurodegenerative disorders. The other two classes of ionotropic glutamate receptors (AMPA and kainate) appear less relevant in HAND and will not be discussed in further detail in this review. NMDAR are heterotetrameric structures that are subtyped depending upon the subunit structure of the receptor. There are three NMDAR subunit groups: NMDAR subunit 1 (NR1), NR2, and the more recently described NR3, which appears to have roles in NMDAR regulation. In most receptors one NR1 subunit (of which there are eight splice variants, NR1a-h) is combined with one NR2 subunit (of which there are four (NR2A-D) differing by the gene or origin), forming NR1/NR2 heterodimers, two of which make up a functional NMDAR (Köhr 2006; Guillemin 2005a; Dingledine et al. 1999; Chatterton et al. 2002). Neurons expressing a high proportion of NMDARs which contain the NR2A and NR2B subunits (i.e., hippocampal, forebrain, and basal ganglia neurons) appear particularly sensitive to NMDAR-mediated excitotoxicity and indeed are particularly vulnerable in HIV infection (O'Donnell et al. 2006).

NMDAR activation in HAND is a result of several mechanisms, some of which are unique to HIV infection when compared with other neuroinflammatory and neurodegenerative disorders. Firstly, neuronal NMDAR activation follows elevated extracellular glutamate as a result of both increased presynaptic release and impaired clearance from the synaptic space. Secondly, high levels of QUIN (an NMDAR agonist) released by activated macrophages (and to a lesser extent microglia) lead to NMDAR activation and excitotoxicity. Thirdly, there are the direct effects of some HIV-related proteins which can both activate the NMDAR and augment receptor activation by glutamate and other agonists. Furthermore, astrocytes play an important homeostatic role in protecting cells from excitotoxic cell death by efficient uptake of glutamate and other neurotransmitters from the extracellular environment in healthy and diseased brains (reviewed by Sofroniew and Vinters 2010), a process which is disturbed in HAND and contributes to excitotoxicity.

Importantly, oxidative stress and lipid peroxidation also contribute to HAND and may occur purely because of downstream effects of NMDAR-mediated excitotoxicity, although there are direct effects of cytokines/chemokines or HIV proteins (particularly gp120 and tat) (reviewed by Mattson et al. 2005). The relative significance of these multiple pathways to cell death and dysfunction in HAND is difficult to establish with certainty, particularly as oxidative stress, inflammation, and excitotoxicity share common final pathways to cell death by caspase activation and apoptosis. Furthermore, some of these pathways are sequential (rather than convergent) and therefore pathophysiologically inseparable. Additionally, caspases, p53 and other enzymes, and transduction systems that mediate apoptosis seem also to play a role in neuronal dysfunction, injury, and dendritic degeneration in the absence of cell death (Garden et al. 2002, 2004).

Alternative pathological mechanisms have been recently advanced and may act simultaneously with excitotoxicity to produce HAND. Of note in particular, HIV has been shown to additionally infect neural progenitor cells and adversely impact on adult neurogenesis (ANG) (the process of neuron generation from neural progenitor cells possibly occurring in adults), likely by perturbing neurotrophic support from astrocytes and altered metabolic processes in particular insulin homeostasis in the CNS (reviewed by Lindl et al. 2010; Mattson et al. 2005). Neuronal autophagy (a protective process by which terminally differentiated neurons process and degrade aged proteins within the cytosol ultimately leading to cell death) is also inhibited by HIV-infected microglial cells (Alirezaei et al. 2008a, b). Inhibition of autophagy is associated with the development of neurodegenerative disorders. HIV-induced inflammation also leads to increased levels of ubiquitinated proteins as a result of disruption of the immunoproteasome function, and thus disturbance in neuronal and synaptic function, hypothesized to contribute to neurocognitive impairment (Nguyen et al. 2010).

## 5 Neuropathogenesis of HAND in the Era of HAART

Importantly, there have been relatively few histopathological studies examining HAND-affected individuals, particularly in the pre-AIDS stage of illness (Bell 2004) and in the era of HAART. After the introduction of antiretroviral therapy (but before widespread use of HAART), HIVE remained at high levels, for example, 26 % in one autopsy study (Masliah et al. 2000). Despite its efficacy in the prevention of opportunistic infections, HAART does not fully protect from HIVE, and "burnt-out" forms of HIVE have been observed in autopsy studies in those treated with HAART, displaying neuronal loss and leukoencephalopathy, which correlate with worsening cognitive function prior to death (Gray et al. 2003). An important issue that follows from these observations is whether the clinical disorder of HAND in the current era is pathologically indeed the same disorder as that observed before the widespread use of HAART as several clinical and pathological changes have been observed. While an association between the presence and severity of HAD and HIV proviral load in the brain (Bell et al. 1998) and CSF, or markers of inflammation (such as beta-2 microglobulin), has been historically described, it is not consistently demonstrated in HAART-treated patients (Brew et al. 2007). Use of antiretroviral agents with high CSF penetrance and activity against resting monocyte-lineage cells in the CNS has not been demonstrated in human trials to consistently yield beneficial effects on neuropsychological function,

though this may be related to trial design issues (Cysique et al. 2011; Brew et al. 2007). While further evaluation of this therapeutic approach is required, this implicates processes that are independent of HIV itself as responsible for ongoing HAND neuropathogenesis. In addition, cognitive changes are now not as closely correlated with basal ganglia abnormalities, with changes in the mesial temporal lobe possibly more relevant (reviewed by Brew 2004). Furthermore, HAART itself leads to upregulation of lymphocytes with immune reconstitution (Hirsch et al. 2004), which may alter the neuropathology in HIV-infected individuals on treatment (Bell 2004). Additionally, as the HAART-treated population ages, superimposed or synergistic effects of other comorbidities may also impact on the clinical expression of HAND, particularly vascular dementia, Alzheimer's dementia (Brew 2004), and other medical disorders, or toxicity arising from HAART itself. Therefore, while excitotoxicity in HAND has definitely been demonstrated, clearly ongoing research is required to establish its relative importance currently and with further therapeutic advances.

## 6 Mechanisms of Excitotoxicity in HAND

## 6.1 Disturbed Glutamate Homeostasis

Excessive extracellular glutamate concentrations occur in HAND patients, reflected by increased concentrations of glutamate in the CSF (Ferrarese et al. 2001) and reduced total parietal glutamate (due to reduced intracellular (neuronal) glutamate levels) in HAND-affected individuals (Ernst et al. 2010). Increased extracellular glutamate could occur as a result of impaired clearance of glutamate from the synaptic space and/or increased glutamate release presynaptically.

The process of glutamate transport (and thus clearance) from the synaptic space occurs primarily through the actions of excitatory amino acid transporter 2 (EAAT2) located on pre- and postsynaptic neuronal membranes and astrocytes sheathing synapses in the CNS (Sheldon and Robinson 2007). Reduced astrocyte clearance of glutamate from the extracellular space has been demonstrated in HIV infection, firstly due to impaired transporter function, following dysregulated release of glutamate by astrocytes, and as a consequence of overt astrocyte loss.

Gp120 activates astrocytic Na+/H+ antiporters in rat and human astrocytes in vitro, leading to depletion of intracellular H+, indirect induction of outward K+ movement due to intracellular alkalinization, and a reduction in the Na+ gradient across the astrocytic membrane. These changes are thought to be responsible for reduced astrocyte uptake (and therefore excessive extracellular levels) of glutamate (Patton et al. 2000). Furthermore, gp120 induces transcriptional downmodulation of the EAAT2 transporter gene in both infected and uninfected human astrocytes and attenuates glutamate transport by the cells in vitro (Wang et al. 2004). HIV-infected astrocytes also demonstrate higher levels of astrocyteelevated gene-1 (AEG-1) expression, known to decrease EAAT2 promoter activity (and thus EAAT2 expression) (Sheldon and Robinson 2007 and references therein). In addition to adversely impacting on astrocyte uptake of glutamate, gp120 can stimulate the release of glutamate from astrocytes into the extracellular space (Benos et al. 1994).

QUIN (further detailed below) also impairs the function of EAAT2 on astrocytes and adversely impacts the glutamate-glutamine cycle within human fetal astrocytes (by inhibition of glutamine synthase (GS) enzyme) and therefore likely further contributes to the accumulation of glutamate in the extracellular environment (Ting et al. 2009). There is also evidence from rat models that QUIN impairs the uptake of glutamate (but not inhibitory neurotransmitters) by synaptic vesicles, representing another avenue leading to excessive extracellular glutamate (Tavares et al. 2000).

Loss of astrocytes in HIV infection clearly also can contribute to increased extracellular glutamate and therefore excitotoxicity and more broadly to the loss of trophic support, synaptic homeostasis, CNS metabolism, and maintenance of the blood–brain barrier (reviewed by Sofroniew and Vinters 2010). QUIN induces apoptosis of human astrocytes in vitro at concentrations comparable found in patients with HAND, in a dose-dependent manner (Guillemin et al. 2005b).

Astrocytes also display functional NMDAR that can be activated by QUIN (Braidy et al. 2009; Lee et al. 2010). Furthermore, astrocytes can support restricted HIV infection and sustain production of tat and nef, both of which can independently adversely impact on neuronal function and lead to neuronal death, as detailed below.

## 6.2 Role of the Kynurenine Pathway and Quinolinic Acid in Excitotoxicity

QUIN is an intermediate in the kynurenine pathway (KP), the major route of L-tryptophan catabolism (ending in the production of nicotinamide adenine dinucleotide (NAD)), and is implicated in the pathogenesis of many neuroinflammatory disorders, including HAND. QUIN is predominantly produced in the CNS by macrophages/microglia, after induction of the KP by proinflammatory cytokines, some interferons (particularly IFN- $\gamma$ ), and possibly by direct induction by HIV proteins tat and nef (Smith et al. 2001) produced within the CNS. Furthermore, BBB endothelial cells and pericytes can be activated across an intact BBB by inflammatory signals (IFN- $\gamma$  in particular) from the periphery to produce QUIN precursors. These intermediaries can be further metabolized to QUIN by perivascular macrophages and microglia even in the absence of CNS HIV infection (Owe-Young et al. 2008).

QUIN causes dendritic injury at low concentrations (consistent however with levels found in HAND-affected individuals) and neuronal death at higher concentrations (Smith et al. 2001; Kerr et al. 1998) through activation of NMDAR. Neurons of the hippocampus, striatum, and neocortex are particularly sensitive to QUIN (reviewed in Guillemin et al. 2005a). QUIN is an endogenous competitive

partial agonist of NMDAR containing NR1 and NR2A or NR2B subunits (Priestley et al. 1995), the very neurons that are particularly vulnerable to HIV-induced neurotoxicity (O'Donnell et al. 2006).

Indeed, several lines of evidence support a significant contribution of QUIN to the NMDAR-mediated excitotoxicity that occurs in HAND (reviewed by Kandanearatchi and Brew 2012 and by Guillemin et al. 2005a). CSF QUIN levels are elevated in HAND-affected patients and correlate with the severity of HIVassociated cognitive impairment in some studies (Heyes et al. 1991) but not all (Valle et al. 2004). Furthermore, atrophy of brain regions that are particularly vulnerable to excitotoxic injury (but not regions established to be resistant to excitotoxic injury) correlates with elevations in CSF QUIN concentrations in HIV-infected patients (Heyes et al. 2001).

Animal models have also demonstrated neuronal loss in cell cultures following exposure to QUIN (Whetsell and Schwarcz 1989) and regional susceptibility to effects of QUIN in the brain that parallel those regions preferentially affected in HAND (reviewed in Guillemin et al. 2005a). Furthermore, pharmacological blockade of NMDAR activation and the KP upstream of QUIN production (Kerr et al. 1997) have been shown to attenuate the neuronal damage and loss by macrophage supernatants.

HAART appears to reduce CSF levels of QUIN in treated individuals, and mathematical modelling indicates that QUIN decay patterns parallel CSF HIV decrement and improvement in neurological performance (Valle et al. 2004). QUIN also directly activates astrocytes, thus amplifying the inflammatory response and astrogliosis in neuroinflammatory disorders including HAND (Ting et al. 2009 and reviewed in Guillemin et al. 2005a).

#### 6.3 HIV Proteins as Excitotoxins

HIV proteins can adversely impact all resident cells in the CNS (including neurons) and blood-derived cells without requiring direct infection. Defective virions and viral particles, viral particles released by cell death, together with viral coat proteins shed by HIV upon cellular infection, are present in the extracellular environment. HIV-infected cells also express some HIV proteins in their surface and through cell-cell interactions impact on uninfected adjacent cells (Nath 2002). Furthermore, gp120, Tat, and Vpr have all been shown to be present in the serum of HIV-infected patients, and Tat (Schwarze et al. 1999) and gp120 (Annunziata 2003) can cross the (often defective) blood–brain barrier, representing another pathway by which CNS neurons and other cells may be exposed to HIV proteins in the absence of direct infection.

There is evidence that gp120 (and Tat to a lesser extent) acts directly as excitotoxins through NMDAR activation in HAND, further detailed below. Other HIV proteins have also been studied in isolation; however, currently there is a lack of convincing evidence to support a significant contribution to excitotoxicity. Of note, gp41 has been demonstrated to induce the release of glutamate from

astrocytes in vitro (Nath 2002 and references therein). Nef is toxic to human neurons and microglial cells in vitro (Trillo-Pazos et al. 2000) (possibly through adverse impacts on neuronal potassium currents across the neuronal membrane (Werner et al. 1991)). In vivo nef exposure also leads to monocyte/macrophage recruitment, increased production of TNF- $\alpha$  and astrogliosis, and behavioral change in rats (Mordelet et al. 2004). Vpr has not been established as a contributor to excitotoxicity; however, it has been shown experimentally to induce neuronal apoptosis possibly through activation of caspase-8 (Patel et al. 2000) and neuronal lysis through disturbance of membrane permeability with subsequent depolarization (Piller et al. 1998, 1999). Rev, Vpu, and other HIV proteins have been studied much less than the above proteins, and at this stage there is insufficient evidence describing a direct role of these proteins in the pathogenesis of HAND.

## 6.3.1 Gp120

As part of the HIV life cycle, gp120 forms virion spikes and interacts with receptors on target cells (Reitz and Gallo 2009). In addition, gp120 induces neuronal cell death of midbrain dopaminergic, hippocampal, cortical, and cerebellar granule neurons in vitro and within rodent brains in vivo (Bachis et al. 2003 and references therein). While experimental evidence suggests that gp120 has the capacity to induce neuronal apoptosis through cytokine receptor activation directly (reviewed in Gonzalez-Scarano and Martin-Garcia 2005) and involvement of the caspase system of enzymes (Garden et al. 2002), the relative importance of this effect pathophysiologically is not certain.

The bulk of evidence suggests that gp120 contributes to neuronal apoptosis ultimately through NMDA receptor-mediated excitotoxicity; however, again the relative importance of this protein overall in the excitotoxicity in HAND is not established. It has been demonstrated experimentally that HIV requires the presence of non-neuronal cells (microglia and macrocytes) to induce excitotoxicity (Meucci and Miller 1996; Kaul et al. 2001, 2005 and references therein), and gp120 has been implicated in the activation of these cells. Indeed, in vivo studies using transgenic gp120 positive mice have demonstrated that gp120 in the absence of HIV itself is sufficient to induce excitotoxic neuronal injury, reduced dendritic density, neuronal death, and behavioral changes consistent with HAND, effects that are at least partly ameliorated by the NMDAR antagonist treatment (Kaul et al. 2005 and references therein). The requirement of non-neuronal cells, however, for neurotoxicity in HIV implies a microglial cell-derived intermediate at play (QUIN, e.g., has been proposed as at least one of these intermediaries as detailed above as well as arachidonic acid (AA)). AA is released by astrocytes and macrophages in the presence of gp120. AA is known to reduce the reuptake of glutamate and other excitatory amino acids by astrocytes and nerve endings via EAA transporters in vitro. AA can also potentiate neuronal NMDAR activation by glutamate, increasing channel open time and mean open probability after activation (Miller et al. 1992; Lipton et al. 1994). There is also some evidence, however, that gp120 itself directly interacts with the NMDAR at its glycine binding site, potentiating its activation by endogenous agonists (Li et al. 2005; Kaul et al. 2001).

#### 6.3.2 Tat

Tat is actively released extracellularly by HIV-1-infected lymphoid and glial cells and astrocytes and can be taken up by most cell types (including neurons) or interact with cell surfaces, thereby exerting toxicity to both infected and uninfected cells. Tat also appears to evade proteolysis by neurons and enter axonal transport systems and therefore can cause neuronal damage and microglial activation at sites distant from cells infected with HIV (reviewed by Li et al. 2009). Cells particularly sensitive to the neurotoxic effects of Tat are those in the striatum, dentate gyrus, and CA3 region of the hippocampus (Rumbaugh and Nath 2006).

Experimental evidence supports a role of Tat in NMDAR-mediated excitotoxicity by potentiating its activation by endogenous agonists (Haughey et al. 2001; Li et al. 2005; Mattson et al. 2005). The pathway of receptor potentiation involves phosphorylation of NMDAR subunits NR2A and NR2B and modulation of Zn-binding sites on the NMDAR (Haughey et al. 2001). This potentiation may explain the observed synergy between Tat and gp120 (Nath et al. 2000; Ferris et al. 2008), as well as with glutamate itself (Li et al. 2009 and references therein), in inducing excitotoxic neuronal cell death. Furthermore, Tat uniquely activates the NMDAR, at a site distinct from glutamate. In vitro rat hippocampal neurons are activated by Tat binding to a distinct modulatory site on the NR1 subunit of NMDARs at concentrations compatible to those in the brains of HIV-infected individuals (Song et al. 2003). The relative importance of this interaction with the NMDAR is uncertain, however, in the pathogenesis of HAND.

Like gp120, there is extensive evidence of Tat involvement in the widespread glial cell activation characteristic of HIVE (particularly monocytes and macrophages), with subsequent induction of a wide range of cytokines (Nath 2002; Mattson et al. 2005; Rumbaugh and Nath 2006). Indeed, Tat increases NO production by microglial cells through inducible NOS gene transactivation, an effect that is amplified in the presence of IFN- $\gamma$  (Polazzi et al. 1999). Similar to gp120, Tat has also been demonstrated to bind and activate some chemokine receptors directly (Albini et al. 1998). Furthermore, Tat interferes with multiple molecular mechanisms in microglial cells experimentally, potentially further contributing to neuropathology (reviewed by Minghetti et al. 2004). Importantly, however, the relative importance of these multiple pathways to neuronal death induced by Tat experimentally is uncertain. Interestingly, astrocytes transfected with Tat are protected from oxidative stress and cellular injury experimentally, despite the extracellular release of Tat inducing neurite trimming and neuron death (Chauhan et al. 2003).

## 7 Adjunctive Therapeutic Approaches: The Role of NMDAR Antagonists

Current HAART fails to protect HIV-infected individuals from HAND even when viral suppression is achieved and immune system function optimized. Therefore, therapies that augment the partially protective effects of HAART are actively being sought. Multiple avenues for alternative therapies have been explored.

The following discussion will focus on agents that reduce excitotoxicity at the level of NMDAR activation; however, the associated rise in intracellular calcium with activation of NOS and apoptotic pathways and/or the impacts of locally and peripherally produced cytokines are also acknowledged as potential targets (Wallace 2006). For example, both lithium and valproate modulate apoptotic protein function; lithium also stabilizes Na+/K+ channels and reduces inflammation.

Notably, studies have shown preliminary efficacy of valproate in preventing neuronal dysfunction in animal models of HIVE (Dou et al. 2003, 2005) and in the treatment of HAND-affected individuals (Schifitto et al. 2009; Letendre et al. 2006).

The recognition that NMDAR activation plays a central role in neuronal degeneration and death in HAND has been facilitated by the use of NMDAR antagonists experimentally. Therapeutic trials of such antagonists have naturally progressed from these findings, however as yet their effectiveness in improving cognitive function or slowing the progression of HAND has not been established. Additionally, while changes to glutamate receptor functioning in astrocytes and microglial cells have been implicated in excitotoxic cell death cascades, the consequences of therapeutic manipulation of glutamate transporters in models of other neurodegenerative disorders have yielded conflicting results (Sheldon and Robinson 2007) and as yet have not been evaluated in the setting of HAND.

Memantine is a noncompetitive, low-affinity, open-channel blocker of the NMDAR (pharmacology reviewed by Lipton 2004), with preferential action on extrasynaptic NMDAR (those containing NR2B subunits) (Shin et al. 2011). It has an acceptable side effect profile due to its ability to prevent NMDAR activation at pathophysiological levels without compromising normal neuronal function; indeed, many experimental agents have failed in this regard with resultant excessive toxicity (Lipton 2004). It is used widely in the treatment of Alzheimer's dementia; however, as yet therapeutic trials examining its use in HAND have not led to recommendations for its use in this setting thus far. In addition, memantine is likely only to be effective when used preventatively or to slow dementia progression in affected patients, as it would have no impact on neuronal impairment once established (Wallace 2006).

Specifically, in vitro studies have demonstrated that neurons are protected from injury and death induced by gp120 (Lipton 1992) and gp120 and tat cotreatment (Nath et al. 2000) in the presence of memantine. Gp120-induced reductions in dendritic and presynaptic terminal density are also diminished in gp120+ transgenic mouse models when treated with memantine (Toggas et al. 1996). Furthermore, memantine has been demonstrated experimentally to prevent Tat-induced death of rat hippocampal neurons in culture. Interestingly, while failing to protect from Tat-induced synapse loss in this study, when administered after significant synapse loss had occurred, memantine induced a recovery of synapse number. The authors hypothesize that NMDA receptor subtypes differentially contribute to synaptic death and recovery and that synaptic NMDA receptor subtypes undergo pharmacological change during the neurotoxic process (Shin et al. 2011).

Memantine treatment in the setting of mild to severe forms of HAND has so far been demonstrated to be safe and tolerable in a double-blind 16-week trial (followed by 20-week washout period), with proton magnetic resonance spectroscopy (MRS) data (measuring metabolites associated with glia and neuronal function in vivo) across the treatment period consistent with a neuroprotective effect in treated individuals (Schifitto et al. 2007). However, the open-label phase of this same study extended to 60 weeks of therapy yielded no statistically significant improvement in neuropsychological performance in those receiving memantine (Zhao et al. 2010). Importantly, however, study limitations necessitate further evaluation to more accurately assess the efficacy of memantine in those affected by HAND (Zhao et al. 2010).

#### 8 Conclusion

So far current therapy for HIV infection has failed to protect treated patients from the development of HAND despite apparent efficacy in virologic control, as least as measured outside of the brain. While current evidence supports the contribution of multiple pathophysiological pathways in the development of HAND, a central role of NMDAR-mediated excitotoxicity is supported by both direct and indirect evidence. Indeed, HIV impacts on this pathway in some ways that are unique among other neuroinflammatory and neurodegenerative disorders, as outlined. Clearly further research is required in the area of HAND pathogenesis to establish the relative importance of NMDAR-mediated excitotoxicity in the development of these disorders (especially in the era of HAART). This would assist in the development of adjunctive therapies in the prevention and treatment of HAND.

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# Excitotoxicity in the Pathogenesis of Autism

M. M. Essa, Nady Braidy, S. Subash, R. K. Vijayan, and Gilles J. Guillemin

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#### Abstract

Autism is a neurodevelopmental disorder characterized by stereotyped interests and behaviors and abnormalities in verbal and nonverbal communication. Autism is reported as a multifactorial disorder resulting from interactions between genetic, environmental, and immunological factors. Excitotoxicity and oxidative stress are potential mechanisms, which are likely to serve as a converging point to these risk factors. Numerous studies suggest that excitotoxicity is a likely cause of neuronal dysfunction in autistic patients. Glutamate is the main excitatory neurotransmitter generated in the CNS, and

M.M. Essa (🖂) • S. Subash • R.K. Vijayan

N. Braidy

Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

e-mail: n.braidy@unsw.edu.au

G.J. Guillemin St. Vincent's Centre for Applied Medical Research, St. Vincent's Hospital, Sydney, Australia

Neuropharmacology Group, MND and Neurodegenerative Diseases Research Centre, Macquarie University, North Ryde, NSW, Australia e-mail: gilles.guillemin@mq.edu.au

Department of Food Science and Nutrition, College of Agriculture and Marine Sciences, Sultan Qaboos University, Muscat, Oman

Ageing and Dementia Research Group, Sultan Qaboos University, Oman e-mail: drmdessa@gmail.com

overactivation of glutamate receptors triggers excitotoxicity. Hyperactivation of glutamatergic receptors, NMDA and AMPA, leads to activation of enzymes, which damage cellular structure, membrane permeability, and electrochemical gradients. The role of excitotoxicity in autistic subjects is summarized in this chapter.

#### **Keywords**

Autism • Excitotoxicity • Free radicals • Glutamatergic receptors • Ion channel • Membrane potential • Neurotransmitter

#### 1 Introduction

Autism is a neurodevelopmental disorder associated with several cognitive manifestations including repetitive behaviors, social withdrawal, and abnormalities in verbal and nonverbal communications (Gillberg and Coleman 2000). Baron-Cohen et al. (2009) showed that autism affects 1–2 in 100 children, according to recent data on the broad array of autism spectrum disorders (ASD) (Baron-Cohen et al. 2009).

Excitotoxicity is an important pathological event mediated by neuronal excitation through overstimulation of neurons by excitatory amino acids receptors of glutamate and aspartate (Olney 1969). The pathological hallmarks of excitotoxicity include neuronal swelling, vacuolization, and neuronal dystrophy (Farooqui and Horrocks 1994). Under physiological conditions, excitatory receptors induce the influx of sodium, calcium, and potassium, which results in neuronal excitation (Lan et al. 2001). Excess levels of glutamate and other excitatory molecules result in overexcitation of the ionotropic glutamatergic receptors N-methyl-D-aspartate (NMDA) and 2-amino-3-(3-hydroxy-5-methylisoxazol-4-yl) propionate (AMPA) and the release of other excitotoxins followed by excessive calcium levels in cytosol. The influx of intracellular calcium triggers the induction of inducible nitric oxide (iNOS) and phosphorylation of protein kinase C. Increased iNOS enhances nitric oxide (NO•) production in excess, whereas protein kinase C activates phospholipase A2 which in turn results in the generation of pro-inflammatory molecules (Babu et al. 1994; Banion 1999). The subsequent generation of free radicals can inhibit oxidative phosphorylation and damage mitochondrial enzymes involved in the electron transport chain, which mitigate energy production (Eliasson et al. 1999).

Seizures represent important pathological event associated with excitotoxicity in autistic children. Experimental studies substantiated the relationship between seizure formation and excitotoxicity (Rogawski 1995). Elevation of glutamate levels in brain is the primary cause of seizure formation (Johnston 1995; Blaylock 2003).

## 2 Evidence for Excitotoxicity in Autistic Patients

Biochemical analysis has shown elevated serum level of glutamate in autistic patients compared to controls (Shinohe et al. 2006). Abnormalities in immune

system regulation have also been reported in autistic patients (Cohly and Panja 2005). Higher levels of tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) receptor II have been observed in the sera of children with autism spectrum disorders (Chez et al. 2007). Microglial activation can further enhance the release of TNF- $\alpha$ , which increases reactive oxygen and nitrogen species and inhibits the reuptake of glutamate (Hu et al. 2001).

Increased glutamate may also occur as a result of interaction between pathogens with bacterial components, viruses, and lipopolysaccharides (Fontana et al. 1996). Impaired immunity promotes viral persistence in host tissues for a greater period of time and stimulates the release of glutamate and other excitotoxins, leading to increased production of reactive oxygen species, and energy depletion in neurons (Espey et al. 1998). Peroxynitrates and other peroxidation products reduce the function of mitochondrial enzymes by impairing oxidative phosphorylation and inhibiting complex II of the electron transport chain (Fosslier 2001; Novelli et al. 1988). Moreover, lipid peroxidation products can interact with synaptic protein and impair transport of glucose and glutamate to reduce energy production and increase vulnerability to excitotoxicity (Henneberry 1989).

The formation of seizures is associated with excitotoxicity and the accumulation of glutamate (Rogawski 1995). The infant brain is more vulnerable to excitotoxic effects since the developing brain contains more synaptic glutamate receptors than at birth, and that number slowly declines during the ageing process (Johnston 1995). There is substantial evidence that excitotoxicity is responsible for the mirror focus reported in temporal lobe seizures and cognitive decline associated with epilepsy (Ekonomou and Angelatou 1999; Olney et al. 1986). Pathological changes have been observed in the hippocampus after prolonged seizures in several cases of autism. These changes include the progressive loss of hippocampal neurons in the CA1 and CA3, as well as dendritic swelling in the hilus of the fascia dentate (Ekonomou and Angelatou 1999; Olney et al. 1986).

Excitotoxic processes in the brain can also be observed using magnetic resonance imaging (MRI), usually with cortical and subcortical atrophy and progressive ventricular dilation (Baxter et al. 1996; Gospe and Hecht 1998). A magnetic resonance spectroscopy study by Page et al. (2006) demonstrated that individuals with ASD have significantly higher concentrations of glutamate in the amygdale-hippocampal region than healthy age-matched controls (Page et al. 2006). Another study demonstrated increased NMDA receptor content in several cases of temporal lobe epilepsy, suggestive of dentate granule cell hyperexcitability (Mathern et al. 1999). Postmortem studies have suggested that dysfunction in the glutamatergic system in the cerebellum may also be involved in the pathophysiology of autism. Interestingly, a recent study found that the anatomic substrate of the limbic system is considerably smaller in autistic children compared to controls (Saitoh et al. 2001).

Additionally, Zilbovicius et al. (2000) used positron emission tomography to visualize neuronal damage in 75 % of autistic children. These changes were localized to the auditory association area and temporal cortex, similar to the areas of hypoperfusion reported in a majority of cases with infantile spasms

(Zilbovicius et al. 2000). As well, experimental lesions in the temporal lobe of nonhuman primates closely resemble temporal lesions observed in autistic children (Bachevalier 1994).

## 3 Conclusion

Autism is a multifactorial neurodevelopmental disorder characterized by deficits in neurobehavioral and neurological dysfunction. Excitotoxicity is triggered by potential mutation in ion channels and signalling pathways, viral and bacterial pathogens, toxic metals, and free radical generation. Overexpression of glutamate receptors and increased glutamate levels lead to increased calcium influx and oxidative stress and progressive cellular degeneration and cell death. Genetic defect, such as mutation in voltage gated or ligand channels that regulate neuronal excitability, leads to defect in synaptic transmission and excitotoxic condition in autism.

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## Glutamate and Neurodegeneration in the Retina

## Thomas E. Salt

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#### Abstract

L-Glutamate is a neurotransmitter throughout the various levels of retinal neurons (photoreceptors, bipolar cells, ganglion cells) and acts at a range of ionotropic (iGlu) and metabotropic glutamate (mGlu) receptors to exert its physiological roles. A function unique to the retina can be ascribed to mGlu6 receptors, which are expressed only by ON bipolar cells at the sign-inverting synapse formed by photoreceptors. More conventional excitatory sign-conserving synapses utilizing iGlu receptors are formed at other locations in the retina. Over-activation of iGlu receptors (and especially NMDA receptors) is known to lead to excitotoxicity in the retina, as in other areas of the nervous system, and this may be associated with excessive neuronal  $Ca^{2+}$  influx. This excitotoxicity can be prevented or countered by iGlu receptor antagonists, both in vitro and in vivo. Furthermore, in some experimental models of retinal neurodegeneration, e.g., for glaucoma or ischemia, a number of different types of iGluR antagonists have been shown to be effective in a variety of species. In particular, the uncompetitive NMDA antagonist memantine has shown promise

T.E. Salt

Department of Visual Neuroscience, UCL Institute of Ophthalmology, London, UK e-mail: T.Salt@ucl.ac.uk

in experimental studies, but clinical trials to treat glaucoma with this drug have not had a successful outcome, possibly due to the unavailability of suitable markers of disease progression and heterogeneity of the disease in humans. It is thus important that further work is carried out so as to translate preclinical findings on neurodegeneration into successful clinical treatments.

#### Keywords

AMPA receptor • Glaucoma • mGlu receptor • NMDA receptor • Retina

List of Ab	breviations
AMPA	(S)-α-amino-3-hydroxy-5-methyl-4-iso-xazolepropionic acid
EAAT	Excitatory amino acid transporter
L-AP4	(APB) L-2-amino-4-phosphonobutyric acid
mGlu	Metabotropic glutamate
NMDA	N-methyl-D-aspartate
RGC	Retinal ganglion cell

## 1 Introduction

The excitatory amino acid L-glutamate (henceforth referred to as "glutamate") has been known to have toxic effects in the eye and the retina for over five decades, since the early work of Lucas and Newhouse (Lucas and Newhouse 1957) and Kobayashi (Kobayashi 1967), working in mice and rabbits. Since then the role of glutamate as an excitatory neurotransmitter in the central nervous system and retina has been well established, based on the pioneering work of Curtis and Watkins (Curtis and Watkins 1963) and the subsequent development of a pharmacological receptor scheme for the excitatory amino acids (Watkins and Evans 1981; Lodge 2009; Traynelis et al. 2010).

As with several other neurotransmitter receptor systems, the glutamate receptors can be divided into two families: the ionotropic receptors (Traynelis et al. 2010) and the metabotropic receptors (Niswender and Conn 2010). The ionotropic glutamate receptors, commonly referred to as NMDA receptors (NR1, NR2A-D, NR3A, NR3B), AMPA receptors (GluR1-4), and kainate receptors (GluR5, GluR6, KA1, KA2), are tetra-heteromeric ligand-gated cation channels (Lodge 2009; Traynelis et al. 2010) that can have a wide variety of characteristics depending on their subunit composition. In addition, *delta* receptors have been described that appear to form homomeric receptors that are not activated by any known agonist (Traynelis et al. 2010). A standard IUPHAR-approved naming convention has been developed that supersedes the common nomenclature (Table 1) (Traynelis et al. 2010). Eight metabotropic glutamate (mGlu) receptor subtypes (mGlu1-mGlu8) have been characterized to date, all of which can be found in the retina, with mGlu6 having a particularly unique role in the visual ON pathway (Morgans et al. 2010) (see below). The mGlu receptors can be placed into three Groups (I, II, III) on the basis

			Example anta	gonists
Receptor	Common subunit names	IUPHAR subunit names	Competitive	Uncompetitive
AMPA	GluR1, GluRA	GluA1	CNQX	
	GluR2, GluRB	GluA2	DNQX	GYKI-52466
	GluR3, GluRC	GluA3	NBQX	GYKI-53655
	GluR4, GluRD	GluA4	LY293558	
Kainate	GluR5	GluK1	CNQX	GYKI-53655
	GluR6	GluK2	DNQX	
			NBQX	
	GluR7	GluK3	LY382884	
	KA1	GluK4		
	KA2	GluK5		
NMDA	NMDAR1, NR1	GluN1		Mg <sup>2+</sup>
	NMDAR2A, NR2A	GluN2A	D-AP5	MK-801
	NMDAR2B, NR2B	GluN2B	R-CPP	Phencyclidine
	NMDAR2C, NR2C	GluN2C		Ketamine
	NMDAR2D, NR2D	GluN2D		Memantine
	NR3A	GluN3A		
	NR3B	GluN3B		
delta	δ1, GluR delta-1	GluD1		
	δ2, GluR delta-2	GluD2		

 Table 1
 Ionotropic glutamate receptor subunits

Abbreviations:  $AMPA \alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid, CNQX 6-cyano-7-nitroquinoxaline-2,3-dione, D-AP5 D-(-)-2-amino-5-phosphonopentanoic acid, DNQX 6,7-dinitroquinoxaline-2,3-dione, NBQX 2,3-dioxo-6-nitro-1,2,3,4-tetrahydrobenzo[f] quinoxaline-7-sulfonamide, NMDA N-methyl-D-aspartic acid, R-CPP 3-((R)-2-carboxypiperazin-4-yl)-propyl-1-phosphonic acid

of their sequence homology, their pharmacological characteristics, and the types of intracellular transduction cascade to which they may couple in in vitro expression systems (Niswender and Conn 2010). Group I (mGlu1, mGlu5) receptors typically couple to postsynaptic inositol phosphate metabolism, while Group II (mGlu2, mGlu3) and Group III (mGlu4, mGlu6-8) receptors may couple to an inhibitory cyclic-AMP cascade (Table 2). However, it is becoming evident that these receptors may also couple to alternative signalling pathways in a context-dependent manner and this potentially adds complexity to the possible outcomes of mGlu receptor activation (Niswender and Conn 2010).

## 2 Glutamate Receptors in the Retinal Circuitry

Glutamate is found throughout the stages of retinal circuitry, from the photoreceptors which release glutamate tonically in the dark to the retinal ganglion cells (RGCs) that use glutamate as their major transmitter in the retinal output to the brain (Wassle 2004; Lukasiewicz 2009; Slaughter 2010). Photoreceptors release glutamate onto

Table	2 Metabo	Table 2         Metabotropic glutamate receptor groups and subtypes	groups and subtypes					
			Orthosteric agonists	S	Orthosteric antagonists	nists	Allosteric modulators	lulators
		Transduction		Group		Group		
Group	Receptor	Group Receptor mechanism	Subtype selective selective	selective	Subtype selective selective	selective	Positive	Negative
I	mGlu1	mGlu1 IP3/Ca <sup>2+</sup> cascade		DHPG	LY367385	4CPG	Ro67–4853	CPCCOEt JNJ16259685
	mGlu5		CHPG				CDDPB	MTEP, MPEP
I	mGlu2	mGlu2 Inhibitory	LY395756	LY354740		LY341495	BINA	Ro64-5229
	mGlu3 cAMP	cAMP		LY379268	LY395756		LY487379	
		cascade		APDC				
III	mGlu4	mGlu4 Inhibitory	LSP4-2022	L-AP4		MAP4	VU0361737	
	mGlu6 cAMP	cAMP		L-SOP		MPPG		
	mGlu7 cascade	cascade				CPPG		MMPIP, ADX71743
	mGlu8		DCPG				AZ12216052	
Abbre	viations: Al	Abbreviations: APDC 2R,4R-4- aminopyrrolidine-2,4-dicarboxylate, BINA 3'-[[(2-cyclopentyl-2,3-dihydro-6,7-dimethyl]-1-oxo-1H-inden-5-yl)oxy]methyl]-	olidine-2,4-dicarboxy	/late, BINA 3'-[[	(2-cyclopentyl-2,3-	dihydro-6,7-dime	thyl-1-oxo-1H-	inden-5-yl)oxy]methyl]-
(hydrc	1,1 -DIPINENYIJ-4-CATOOXYII hydroxyimino)cyclopropa	1,1 - оприенут]-4-сатоохуис аси, с <i>илив</i> э-суапо-N-(1,5-оприенут-1,1-ругагот-э-удоспласатиас, с <i>иго</i> 2-силот-э-пущохурпенувдусите, с <i>гесиова 1-</i> hydroxyimino)cyclopropa[b]chromen-1a-carboxylate, 4 <i>CPG</i> (S)-4-сатboxyphenylglycine, <i>CPPG alpha</i> -cyclopropy1-4-phosphonophenylglycine, <i>DCPG</i>	arboxylate, 4CPG (	S)-4-carboxyph	or-5-yr)benzamide, enylglycine, <i>CPP</i> C	<i>chru z-c</i> noro-	nyuroxypnen pyl-4-phosphor	ic acid, $CDDFB$ 3-cyano-N-(1,3-cupneny)-111-pyrazot-3-71)oenzamide, $CPFG$ alpha-cyclopropyl-4-phosphonophenylglycine, $DCPG$ [b]chromen-1a-carboxylate, $4CPG$ (S)-4-carboxyphenylglycine, $DCPG$
(S)-3,	4-dicarboxy amino-2-me	(S)-3,4-dicarboxyphenylglycine, <i>DHPG</i> 3,5-dihydroxyphenylglycine, L- <i>AP4</i> L-2-amino-4-phosphonobutyric acid, <i>L-SOP</i> L-serine- <i>O</i> -phosphate, <i>MAP4</i> (S)-2-amino-2-methyl-4-phosphonobutanoic acid, <i>MMPIP</i> 6-(4-methoxyphenyl)-5-methyl-3-(4-pyridinyl)-isoxazolo[4,5-c]pyridin-4(5H)-one, <i>MPEP</i>	,5-dihydroxyphenylg c_acid, MMPIP	Iycine, L- $AP4$ 1 6-(4-methoxyph	L-2-amino-4-phosphenyl)-5-methyl-3-(4	nonobutyric acid 4-pyridinyl)-isox	, L-SOP L-seri azolo[4,5-c]pyr	ne-O-phosphate, MAP4 din-4(5H)-one, MPEP
2-met	hyl-6-(phen	2-methyl-6-(phenylethynyl)-pyridine, MPPG (RS)-α-methyl-4-phosphonophenylglycine, MTEP 3-((2-methyl-1,3-thiazol-4-yl)ethynyl)pyridine.	G (RS)-α-methyl-4-μ	ohosphonopheny	'lglycine, MTEP 3-(	((2-methyl-1,3-th	iazol-4-yl)ethyı	ıyl)pyridine.

either ON or OFF bipolar cells, and it is at this point in the retinal circuitry that opposite polarity light responses are first generated from the "OFF" response of the photoreceptors which reduce their glutamate release upon light stimulation. The ON bipolar cells express mGlu6 receptors in a highly specific manner (indeed this is almost the only location in the nervous system where this receptor is expressed to any significant extent (Nakajima et al. 1993)), and it has been shown that mGlu6 is an essential component of the retinal ON pathway (Nakajima et al. 1993; Morgans et al. 2010). In the dark, constant release of glutamate from photoreceptors activates mGlu6 receptors, and conversely light stimulation leads to a reduction of mGlu6 receptor activation due to a drop in glutamate release onto bipolar cells. This lightinduced deactivation of the mGlu6 signal transduction pathway results in the opening of a nonselective cation channel that is now known to be TRPM1 (Morgans et al. 2010), and this generates the ON response of the ON bipolar cells, representing a sign inversion of the synaptic input from the photoreceptors, mGlu6 receptors can be activated by the synthetic glutamate analogue L-2-amino-4-phosphonobutyric acid (L-AP4, formerly known as APB), and application of this agonist to the retina functionally blocks the ON pathway as it achieves constant activation of mGlu6 receptors, mimicking the constant release of glutamate in darkness (Slaughter 2010). By contrast, OFF bipolar cells express AMPA and kainate receptors rather than mGlu6 receptors and this results in a synaptic OFF response that follows the polarity of the photoreceptor response (Lukasiewicz 2009; Slaughter 2010).

Retinal bipolar cells are predominantly glutamatergic, although glycinergic bipolar cells have also been described (Wassle 2004). NMDA receptors appear in the retinal circuitry postsynaptic to ON bipolar cells and OFF bipolar cells on RGCs and amacrine cells where they participate in both ON and OFF signalling pathways together with AMPA and kainate receptors (Lukasiewicz 2009; Slaughter 2010). There is spatial segregation of ON bipolar cell synapses and OFF bipolar cell synapses onto ON and OFF RGCs, respectively, and NMDA receptors appear to be localized primarily peri-synaptically at ON synapses, whereas they have a more subsynaptic location at ON synapses. Furthermore, it appears that different NMDA receptor subunit types are found at these locations, with NR2B-containing receptors more prevalent at ON synapses and NR2A-containing receptors more prevalent at OFF synapses of RGCs (Fletcher et al. 2000; Zhang and Diamond 2009). In addition to these complexities, the amino acids glycine and D-serine act as co-agonists at NMDA receptors, particularly those containing the NR2B subunit (Traynelis et al. 2010), and it is possible that these co-agonists have a specific role in ON pathway signalling (Gustafson et al. 2007; Kalbaugh et al. 2009).

In addition to the highly specific role of mGlu6 receptors in the ON pathway (see above), the other mGlu receptors are also found in the retina, but specific functional roles have not been so readily ascribed (Koulen et al. 1996, 1997; Dyka et al. 2004; Quraishi et al. 2007). However, a role for mGlu8 as a presynaptic receptor located on photoreceptors and bipolar cells is beginning to emerge, and it appears that at both of these locations, it may function as an autoreceptor to regulate the release of glutamate from photoreceptors and from bipolar cells (Koulen and Brandstatter 2002; Quraishi et al. 2010).

#### 3 Glutamate Receptors in Neurodegenerative Mechanisms

Since the initial observations on retinotoxic effects of glutamate by Lucas and Newhouse (1957), there have been a vast number of studies in many species that have investigated the neurotoxic effects of glutamate and glutamate receptor agonists and neuroprotective effects of antagonists and modulators throughout the nervous system, and these have been extensively reviewed (Choi 1988; Danysz and Parsons 1998; Doble 1999; Waxman and Lynch 2005; Casson 2006; Luo et al. 2011). There has been a substantial focus on NMDA receptors, partly due to the early pharmacological characterization of NMDA receptors (Watkins and Evans 1981) but also due to the observation that neurotoxicity was caused by large amount of Ca<sup>2+</sup> influx induced by over-activation of NMDA receptors, a mechanism that was termed *excitotoxicity* (Choi 1988; Luo et al. 2011). More recently, it has become evident that the subunit composition of NMDA receptors is also critical in how they may take part in excitotoxic processes, and in particular, incorporation of NR3 subunits may have a neuroprotective effect (Nakanishi et al. 2009; Low and Wee 2010; Luo et al. 2011).

It should also be noted that activation of kainate or AMPA receptors can also lead to excitotoxicity (Choi 1992). In particular, the presence of GluR2 in AMPA receptors results in low  $Ca^{2+}$  permeability in most situations, due to RNA editing of the GluR2 subunit at the so-called Q/R site. Thus, GluR2 content and Q/R editing is critical in controlling the  $Ca^{2+}$  permeability of the AMPA receptor, and it has been suggested that increased proportion of unedited GluR2 and reduction in the total amount of overall GluR2 in AMPA receptors makes neurons more susceptible to AMPA-receptor-mediated excitotoxicity (Kim et al. 2001; Lebrun-Julien et al. 2009).

## 4 Glutamate Transporters in Neurodegenerative Mechanisms

It has become evident that, in addition to glutamate receptors, the uptake of glutamate by neurons and glia via specific transporters (Excitatory Amino Acid Transporters, or EAATs) is a key component of synaptic transmission and excitotoxic processes, maintaining a physiological extracellular glutamate level and thus regulating the balance between signalling and pathology (Shigeri et al. 2004; Beart and O'Shea 2007; Bringmann et al. 2009; Furuya et al. 2012). In addition, it has also been suggested that EAATs provide glutathione precursors such as glutamate and cysteine to cells and that this is crucial to prevent oxidative stress and cell death in neurons (Harada et al. 2007; Had-Aissouni 2012). There are five major subtypes of EAAT that have been termed EAAT1-5 in humans with homologues in other mammals and vertebrates (Beart and O'Shea 2007). Within the retina, EAAT1 (also known as GLAST) is localized in Müller cells (the major retinal glial cells) and astrocytes (Rauen et al. 1997). EAAT2 (also known as GLT-1) is mainly localized in cone photoreceptors, cone bipolar cells, RGCs, and Müller

cells (Rauen et al. 1997; Furuya et al. 2012). EAAT3 (also known as EAAC1) is found on Müller cells and neurons including RGCs (Kugler and Beyer 2003; Furuya et al. 2012). EAAT4 has been found in photoreceptors and astrocytes albeit at lower levels than in the brain (Ward et al. 2004; Pignataro et al. 2005). EAAT5 is present in rod photoreceptors and bipolar cells of the retina (Fyk-Kolodziej et al. 2004; Lee et al. 2012), where it may serve to regulate glutamate levels and also modulate a chloride conductance in bipolar cells (Wersinger et al. 2006).

#### 5 Glutamate Excitotoxicity in Retinal Diseases?

The death of RGCs and other retinal neurons upon injection of glutamate agonists has been reported in numerous studies conducted in many vertebrate species (Baltmr et al. 2010; Almasieh et al. 2012). Indeed use of NMDA to lesion RGCs is often used as a model of retinal diseases such as glaucoma (Baltmr et al. 2010; Almasieh et al. 2012). The question thus arises as to whether these excitotoxic mechanisms can be related to retinal disease processes that may occur in, for example, glaucoma or retinal ischemia. There are several strands of evidence to suggest that this may be the case. The most compelling of these is that in in vivo experimental models of glaucoma or ischemia produced by a variety of different means in different species from rodent to nonhuman primate, a range of NMDA receptor antagonists (administered locally or systemically), including MK-801 (Chaudhary et al. 1998; Schuettauf et al. 2000; Guo et al. 2006), memantine (Kim et al. 2002; WoldeMussie et al. 2002; Hare et al. 2004), ifenprodil (Guo et al. 2006), and eliprodil (Kapin et al. 1999), have proven to be effective. Similarly AMPA/kainate antagonists have also been shown to be effective (Schuettauf et al. 2000). Furthermore, genetic deletion of EAATs in mice produces an RGC pathology that bears similarities to glaucoma (Harada et al. 2007).

It has also been reported that glutamate levels are elevated in the vitreous of the eye in glaucoma (Dreyer et al. 1996; Brooks et al. 1997), although these findings have been called into question (Dalton 2001), and contradictory results have been published (Carter-Dawson et al. 2002; Kwon et al. 2005). Although superficially it might be attractive to think that glutamate could be elevated in the vitreous in glaucoma, it would seem that this would be unlikely given the prevalence of uptake and compartmentalization of glutamate in neural tissue (Meldrum 2000; Salt and Cordeiro 2006; Almasieh et al. 2012). Thus, a lack of detectable glutamate increase in the vitreous of the eye should not be taken to indicate that glutamate is not taking part in neurodegenerative processes (Salt and Cordeiro 2006; Osborne 2009).

Memantine is an uncompetitive (open channel blocker) NMDA antagonist with fast kinetics that is in clinical use for neurodegenerative conditions such as Alzheimer's disease (AD) in some countries (Lipton 1993; Parsons et al. 1999; Lipton 2004). This tolerability and apparent efficacy has made it an attractive candidate for retinal neuroprotection in glaucoma (Greenfield et al. 2005; Casson 2006). However, although preclinical data in experimental glaucoma provided positive data for memantine (Kim et al. 2002; WoldeMussie et al. 2002;

Hare et al. 2004; Yucel et al. 2006), positive data from clinical trials have not been forthcoming (Allergan Inc 2008; Osborne 2009). Although disappointing, this result could be due to a number of factors. For example, in addition to NMDA receptors, other factors are likely to contribute to glaucoma (Weinreb and Khaw 2004; Osborne 2009; Almasieh et al. 2012), and blockade of NMDA receptors may not be appropriate or sufficient in every case. Furthermore, it is possible, if not likely, that the end points and methods for monitoring disease progression and benefit of therapy were not appropriate (Osborne 2009). In this respect, recent advances in imaging of different phases of retinal neurodegeneration may be of benefit in any future clinical trials (Cordeiro et al. 2010).

### 6 Neuroprotection via Activation of Receptors

All of the known mGlu receptors have been found in the retina (Hartveit et al. 1995; Koulen et al. 1996, 1997; Dyka et al. 2004; Quraishi et al. 2007). It is known that activation of the Group II or the Group III mGlu receptors can be neuroprotective in some experimental situations (Copani et al. 1995; Bruno et al. 2001). In particular, systemic application of Group II-selective agonists is neuroprotective in wholebrain ischemia models (Bond et al. 1998); in the retina, the Group II agonist LY354740 has been shown to be neuroprotective against RGC apoptosis in a rat glaucoma model (Guo et al. 2006) and against anoxic damage in the goldfish (Beraudi et al. 2007). The mechanism of action remains to be elucidated, but it is known from work in cortical cells that activation of mGlu3 receptors on astrocytes rather than mGlu2 receptors on neurons is needed for protection against NMDAinduced excitotoxicity (Corti et al. 2007), and it may be that this stimulates production of certain growth factors (e.g., GDNF, TGF-beta) (Battaglia et al. 2009; Caraci et al. 2009) which may be important in preventing apoptotic cell death (Almasieh et al. 2012). Although activation of Group III receptors has been suggested to be neuroprotective in the brain (Copani et al. 1995; Bruno et al. 2001; Ngomba et al. 2011), it is unclear whether this would be appropriate in the retina given the importance of the closely related mGlu6 (and mGlu8) receptor in visual processing (see above). In this respect it is noteworthy that mutations of the mGlu6 receptor encoding gene, GRM6, cause visual deficits including night blindness that are attributable to dysfunction of the ON pathway (Dryja et al. 2005; Zeitz et al. 2005, 2007). Thus, Group III mGlu receptors may not be suitable targets for retinal neuroprotection.

## 7 Conclusion

Glutamate receptors play a pivotal role in the normal signalling functions of the retina, as they do elsewhere in the nervous system, with a notable specialization in the function of mGlu6 in the ON pathway. Over-activation of the ionotropic receptors causes excitotoxic neuronal death, and this may be of relevance in the

pathophysiology of retinal diseases such as glaucoma. Neuroprotective strategies aimed at blocking these processes may be useful, but further work is required to translate experimental results to clinical usage. It may be the case that different mechanisms or combinations of mechanisms are applicable in different diseases/ patients, or even at different stages of the same disease. Thus, neuroprotective strategies might need to be tailored to individual patients with appropriate monitoring of disease progression, and this should be attainable in the future (Cordeiro et al. 2010).

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# **Glutamate in the Pathogenesis of Gliomas**

Nady Braidy, Anne Poljak, Tharusha Jayasena, Seray Adams, and Perminder Sachdev

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N. Braidy (🖂) • T. Jayasena

A. Poljak

S. Adams

e-mail: seray.adams@student.unsw.edu.au

P. Sachdev

Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

e-mail: n.braidy@unsw.edu.au; t.jayasena@student.unsw.edu.au

School of Medical Sciences, Faculty of Medicine and Bioanalytical Mass Spectrometry Facility, University of New South Wales, Sydney, Australia e-mail: a.poljak@unsw.edu.au

School of Medical Sciences, Faculty of Medicine, University of New South Wales, Sydney, Australia

Centre for Healthy Brain Ageing, School of Psychiatry, University of New South Wales, Sydney, NSW, Australia

Neuropsychiatric Institute, Prince of Wales Hospital, Sydney, Australia e-mail: P.Sachdev@unsw.edu.au

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#### Abstract

Recent studies have shown that glutamate may serve important roles in the pathobiology of primary brain tumors. Glutamate is produced and secreted from gliomas via specific glutamate transporters as a by-product of glutathione synthesis. Glutamate also plays a major role in the phenotype of malignant gliomas by several mechanisms. The consequential interaction of glutamate with peritumoral neuronal glutamate receptors leads to the development of seizures and excitotoxicity. The latter is thought to promote the expansion gliomas in the vacated surrounding tissue. Glutaminergic receptors such as  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) which lack the GluR2 subunit can activate the AKT and MAPK pathway, to promote the invasion of glioma cells into normal brain tissue. Stimulation of glutaminergic receptors also leads to the activation of focal adhesion kinase, which regulates the proliferation and motility of malignant glioma cells. Given the extensive involvement of glutamate in the development of gliomas, it is likely that pharmacological therapies targeting glutaminergic receptors and glutamate transport may be useful to inhibit glutamate-mediated glioma growth.

#### Keywords

AMPA • Excitotoxicity • Gliomas • Glutamate • Glutathione

#### 1 Introduction

Over the last decade, numerous studies have identified several unique factors that can promote the growth of malignant gliomas. These include a high abundance of vasculature, nutrients, and growth factors that promote the proliferation of tumor cells and specific white matter tracts that pave the way for invading cells to transverse across the brain. Similar to other cancers, the growth of malignant gliomas is dependent on important signalling pathways by activated tyrosine kinases and absence of important tumor suppressor protein such as p53. However, several clinical trials using therapy targeting traditional growth factor pathways such as epidermal growth factor receptor (EGFR), platelet derived growth factor receptor (PDGFR), mTOR, and protein kinase C (PKC) have been unsuccessful (Ohgaki and Kleihues 2009). This suggests that gliomas may possess yet unidentified characteristics which are likely to be of greater therapeutic value.

The excitotoxic neurotransmitter, glutamate, is of major importance in the development of gliomas since (1) it is a very abundant growth factor and (2) it acts as an important motogen in the CNS (Noch and Khalili 2009). Glutamate is released by glioblastomas via system  $x_c^-$  to promote their malignant behavior triggering apoptosis of surrounding tissue and enhancing the invasive nature of gliomas (Ye et al. 1999; Kim et al. 2001). Current experience indicates that 80 % of glioblastomas recurrences occur within 2–3 cm of the perimeter of the original resection cavity (Kim et al. 2009). Therefore, reducing glutamate levels may be empirical for inhibiting the growth of these tumors (Ye and Sontheimer 1999a). Successful treatment of glioma cells may require an increased understanding of glutamate dynamics in the brain to develop novel approaches aimed at improving current ineffective therapies (Savaskan et al. 2011; Simon and von Lehe 2011).

# 2 Glutaminergic System in the CNS

Two broad categories of excitatory amino acid receptors have been identified in the brain: ionotropic receptors which contain an integral ion channel and metabotropic receptors which are coupled to G proteins (Lalo et al. 2011). The NMDA receptor of which there are also subtypes is one of the three types of ionotropic receptors, the other two being  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) and the kainate receptor (Ekici et al. 2012; Fuchs et al. 2012). Glutamate is the primary endogenous excitatory neurotransmitter in the CNS which will activate these receptors. All neurons in the CNS appear to be sensitive to NMDA-mediated toxicity (Savaskan et al. 2011; Simon and von Lehe 2011).

Of the ionotropic receptor channels, the NMDA receptor is unusual in that it is gated by both ligands and voltage (Singh and Kaur 2009). Glutamate is unable to activate the NMDA receptor at normal resting membrane potential (-70 mV) due to the blockage of the ion channel by Mg<sup>2+</sup> ions. Mg<sup>2+</sup> is only removed when the target cell is partially depolarized (e.g., by additional synaptic inputs) to around -30 mV to -20 mV, allowing agonist induced currents to occur. Once activated, the NMDA receptor is permeable to Ca<sup>2+</sup>, Na<sup>+</sup>, and K<sup>+</sup> ions. Ca<sup>2+</sup> entry is thought to activate Ca<sup>2+</sup>-dependent secondary messenger cascade which contributes to certain forms of long-lasting synaptic modification (Frandsen et al. 1989; Mattson et al. 1991). It is thought that sustained increases in intracellular free Ca<sup>2+</sup> due to excessive NMDA receptor activation may be central to neuronal cell death in several neurodegenerative diseases of the brain.

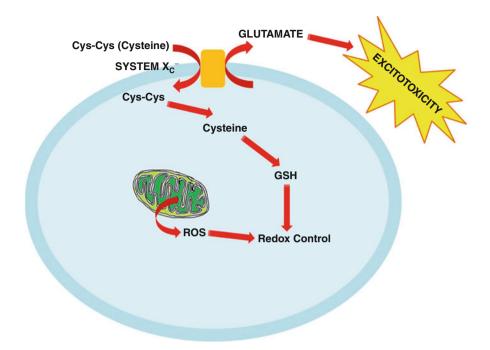
Glutamate uptake is the major route of degradation of glutamate from the extracellular space, making it an important function for nonmalignant astrocytes. The uptake of glutamate into astrocytes occurs predominantly by two Na<sup>+</sup>-dependent excitatory amino acid transporters EAAT1 or EAAT2, otherwise known as Glt-1 and Glast. These transporters are highly localized in the astrocytic membrane (Allritz et al. 2010; Unger et al. 2012). Upon uptake into astrocytes, glutamate is converted to the glutamine by the catalytic activity of glutami synthetase (GS). Glutamine is then released and retaken into neurons where it is hydrolyzed by glutaminases to regenerate glutamate which is stored in synaptic vesicles (Chretien et al. 2002).

# 3 Glutamate Release from Gliomas

In neuronal cells, glutamate is generally released from specific synaptic vesicles in a Ca<sup>2+</sup>-dependent pathway. However, in astrocytes, glutamate is released via a number of different mechanisms including vesicular release (Montana et al. 2006; Ponzio et al. 2006), reverse operation of Na<sup>+</sup>-dependent glutamate transporters (Courtney et al. 1990), swelling of activated anion channels (Kimelberg et al. 1990a, b, c; Kimelberg and Kettenmann 1990), and through specific hemichannels (Ye et al. 2003). Recent studies have failed to associate these pathways as the mechanism of glutamate release in gliomas. These studies suggest that glutamate release in malignant gliomas may occur predominantly via the Na<sup>+</sup>-independent system x<sub>c</sub><sup>-</sup> cysteine glutamate exchanger (Ye and Sontheimer 1999b) (Fig. 1). This important amino acid transporter fosters the release of glutamate from glioma cells concomitantly with the uptake of cysteine necessary for the production of the endogenous antioxidant, glutathione (GSH) (Sato et al. 1999). In primary cells, GSH, which contains the highly reactive -SH group, maintains the normal cellular redox status by binding with reactive oxygen species and nitrosylative radicals. In gliomas, GSH provides a mechanism of protection for the malignant cells by enhancing the cell's redox status and stimulating excitotoxicity in the peritumoral brain (Kandil et al. 2010). Several analogs of the transported amino acids have been shown to disrupt system  $x_c^-$  and impair glutamate release from glioma in vivo (Patel et al. 2004). Additionally, inhibition of system  $x_c^-$  using the inhibitor, L-alphaaminoadipate, leads to more than 50 % reduction in intracellular GSH levels in C6 gliomas, suggesting that the majority of GSH formed in these cells is due to system x<sub>c</sub><sup>-</sup>-mediated uptake of glutamate. Reduced GSH levels due to blockade of system  $x_c^-$  can trigger apoptosis of glioma cells via activation of the JNK and p38 MAPK pathways (Kandil et al. 2010).

Apart from its role in the production of GSH in glioma cells, glutamate can also be converted to  $\alpha$ -ketoglutaraldehyde via the catalytic activity of glutamate dehydrogenase (GDH), which is highly expressed in glioma cells (Yang et al. 2009). Glutamate also serves as an important intermediate for energy production in the tricarboxylic cycle (TCA) due to its metabolism of the essential substrate  $\alpha$ -ketoglutaraldehyde (Yang et al. 2009). Therefore, inhibition of glutamate release from gliomas provides a novel target to reduce the survival of malignant glioma cells by depleting important sources of energy for these glycolysis-impaired cells and reducing their redox defense status.

As previously mentioned, EAAT1–5 have been implicated in glutamate release in both astrocytes and neurons. However, these transporters appear to be dysfunctional, or not expressed in most gliomas (de Groot et al. 2005). One study showed that EAAT2 is highly expressed in low-grade astroglial tumors, but at significantly lower levels in high-grade glioblastoma tumors. Interestingly, the expression of EAAT2 can regulate glioma cell viability. For instance, the presence of EAAT2 in high-grade glioblastoma tumors makes these cells highly vulnerable to cytotoxicity



**Fig. 1** Glutamate is released from glioma cells via system  $x_c^-$  involving the uptake of cysteine necessary for the production of the endogenous antioxidant glutathione (GSH)

via an apoptotic mechanism, and increased EAAT2 expression *ex vivo* can reduce tumorigenicity (de Groot et al. 2005).

Astrocyte elevated gene-1 (AEG-1) is an oncogene that is overexpressed in several tumor cells, including glioblastoma tumors, and is involved in glioma progression (Kang et al. 2005; Lee et al. 2011). AEG-1 has been shown to increase in astrocytes progressing into malignant gliomas, parallel to a decrease in EAAT2 expression (Hu et al. 2009; Baumann and Dorsey 2011; Noch et al. 2011). Recently, Lee et al (2011) showed that AEG-1 repressed EAAT2 expression at the transcriptional level, causing a reduction in glutamate uptake by glial cells, culminating in neuronal cell death (Lee et al. 2011). These findings have also been reiterated in samples of glioma cells obtained from glioma patients indicating a decline in NeuN expression correlating with increased AEG-1 expression (Lee et al. 2011).

#### 4 Effect of Released Glutamate on Neighboring Brain Cells

In the brain, neurons, astrocytes, and microglial and vascular cells are located in close proximity to gliomas, all of which are equally responsive to the released glutamate by a number of different mechanisms.

#### 4.1 Neuron

As earlier discussed, increased levels of glutamate in the extracellular spaces can lead to neurodegeneration by promoting glutamate excitotoxicity. This represents an alternative mechanism for the promotion and expansion of tumoral growth in the adjacent brain tissue. Excitotoxicity is a biochemical process that occurs in response to overactivation of neuronal NMDA receptors in the peritumoral brain, leading to hyperexcitability and seizures which occur frequently in conjunction with gliomas (Moots et al. 1995). Ye et al. (1999) showed that glioma cells release excitotoxic concentrations of glutamate which can induce widespread excitotoxicity in surrounding neurons as well as cellular edema (Ye and Sontheimer 1999a). The loss of these neuronal cells can be prevented by treating neurons with the NMDA receptor antagonist, MK-801/D(-)-2-amino-5-phosphonopentanoic acid, or inhibition of glutamate release from system  $x_c^-$  both in vitro and in tumor-bearing mice (Chung et al. 2005).

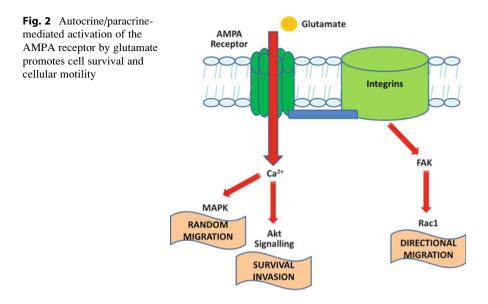
# 4.2 Astrocyte

Unlike neurons and oligodendrocytes, astrocytes are less vulnerable to millimolar concentrations of glutamate (Oka et al. 1993). Astrocytes typically respond to increased neuronal glutamate release by the rapid uptake of glutamate into the cytoplasm using the EAAT1 and EAAT2 transporters. It is likely that the function of these excitatory amino acid transporters may be impaired in astrocytes in the presence of gliomas, or they may lose their ability to catabolize glutamate to the nontoxic metabolite glutamine. Studies have shown that reactive gliosis, which is common during excitotoxicity, induces a loss in GS activity (Ortinski et al. 2010). Such loss may enhance the vulnerability of neurons to glutamate-mediated excitotoxicity and impair the normal homeostatic role of astrocytes in the brain. As well, reduced GS activity can lower GABAergic signalling by disrupting glutamate-glutamine cycling (Ortinski et al. 2010).

Astrocytes also express glutaminergic receptors. NMDA and AMPA ionotropic receptors are functionally expressed in astrocytes (Lee et al. 2010). Overactivation of the NMDA receptor can trigger glial cell death, whereas AMPA activation can alter astrocyte morphology, promote glutamate release from astrocytes, and induce deleterious changes in gene expression and normal cellular function (Weber et al. 2001). Metabotropic glutaminergic receptors may also be altered in response to changes in extracellular levels of glutamate and can modulate the expression of glutamate transporters (Aronica et al. 2003).

# 4.3 Glioma

Ionotropic NMDA, AMPA, and metabotropic glutaminergic receptors are all expressed in glioma cells in cell line cultures and patient-derived glioblastoma



specimens. The NMDA subunits NR2A, NR2B, and NR2C are expressed in several glioma cell lines (Stepulak et al. 2009). Additional studies have demonstrated antitumoral effects with NMDA receptor antagonists (Rzeski et al. 2001). Moreover, kainite receptors have also recently been discovered in several glioma cell lines cultures (Albasanz et al. 1997; Liu et al. 1997).

The AMPA receptor containing the subunits GluR1, GluR3, and GluR4 has been shown to be highly expressed in high-grade gliomas (Ishiuchi et al. 2007). The GluR2 subunit, which is lacking in most gliomas, modulates  $Ca^{2+}$  permeability. If an unedited form of the GluR2 subunit is present,  $Ca^{2+}$  permeable receptors are established. However, the presence of an edited form of the GluR2 subunit makes the AMPA receptor impermeable to  $Ca^{2+}$  and is the case at the synapses. Hence, the absence of the GluR2 subunit in gliomas induces the influx of  $Ca^{2+}$  through the AMPA receptor (Ishiuchi et al. 2007). Furthermore, one study showed that loss of GRIA2, the gene coding for GluR2 expression, leads to a poor prognosis in glioblastomas (Colman et al. 2010). Taken together, this suggests that the AMPA receptor is crucial for the development of glioblastomas. Increased  $Ca^{2+}$  appears to be an important mediator of glioma cell motility and tumor invasion.

Continued release of glutamate can lead to activation of AMPA receptor in the glioma and surrounding cells. Autocrine and paracrine activation of the AMPA receptor has been demonstrated in vivo (Lyons et al. 2007) (Fig. 2). Overexpression of the GluR1 subunit in the AMPA receptor can activate FAK-dependent signalling pathway independent of the AMPA receptor activation (Piao et al. 2009). Although the mechanism associated with FAK activation remains unclear, it may be associated with FERM protein or integrin activity. Activation of the AMPA receptor due to glutamate enhances cellular detachment via Rac1 and stimulates transwell migration in vitro and promotes tumor invasion in vivo (Piao et al. 2009).

Metabotropic receptors, including mGluR3 and mGluR5, have been described in several low- and high-grade glioma cell lines (Caruso et al. 1997; Condorelli et al. 1997). However, characterization of these receptors in human glioma samples remains to be accomplished.

# 5 Targeting the Glutaminergic System as Antitumoral Agents

Given the importance of glutamate in the biology of glioma growth and invasion, current agents which modulate the glutaminergic system may represent potential therapeutic agents. If modulators of the glutaminergic system can limit tumor invasion into the peritumoral brain, these drugs may represent novel therapies which can inhibit glioma infiltration in humans. Several clinical trials have been performed and are summarized here.

# 5.1 Antagonism of the AMPA Receptor

It is well established that glutamate released from gliomas can activate glutaminergic receptors on the same or neighboring cell. However, these drugs are limited due to severe side effects when translated into the clinic (Lipton 2004). Talampanel is a noncompetitive antagonist of the AMPA receptor that can pene-trate the brain and demonstrated a favorable outcome in preclinical studies (Calabrese et al. 2007). The drug has demonstrated an excellent safety profile in a single-arm Phase II study in 32 patients with recurrent glioblastomas or anaplastic gliomas as a monotherapy (Iwamoto et al. 2010). In another larger Phase II study, talampanel was given in newly diagnosed glioblastoma patients in combination with radiation therapy and temozolomide (Grossman et al. 2009). The data collected so far indicates a favorable survival rate with no added toxicity when used in combination. However, additional randomized placebo-controlled trials are necessary to adequately establish the efficacy of this new drug compared with radiation and standard treatment regimes.

# 5.2 Antagonism of Other Glutaminergic Receptors

Other glutamate receptor antagonists may also be of clinical benefit for the treatment of patients with gliomas. Memantine, a noncompetitive NMDA ion channel blocker, is currently approved for the treatment of mild to moderate Alzheimer's disease (AD). Interestingly, preclinical studies have reported reduced glioma proliferation following treatment with memantine (Rzeski et al. 2001). Apart from its antitumoral activity, memantine may also attenuate excitotoxicity in neuronal cells due to increased glutamate in the extracellular space. This may also be important in inhibiting metabolic pathways utilized by malignant glioma cells. Additionally, metabotropic glutamate receptors may also represent a useful target to attenuate NMDA receptor-mediated excitotoxicity and reduce neurological dysfunction, seizures, and increased tumor invasion. LY341495, a selective Group II metabotropic glutamate receptor antagonist, has been shown to reduce glioma proliferation, and clinical trials are under way (Arcella et al. 2005).

# 5.3 Blockade of System x<sub>c</sub><sup>-</sup> in Gliomas

As previously discussed, the system  $x_c^-$  represents the main mechanism of glutamate release from gliomas. This unique transporter can be inhibited using sulfasalazine and FDA-approved drug that can significantly slow down tumor growth in mice (Chung et al. 2005). A Phase I study of this agent in 9 heavily pretreated patients with poor performance was terminated by the investigator due to poor efficacy outcomes.

#### 5.4 Inhibition of Glutamate Reuptake

Enhancing glutamate uptake in glial cells represents an alternative means to prevent glutamate accumulation in peritumoral tissue and reduce glioma invasion in surrounding brain tissue. This notion has been demonstrated in vitro and in preclinical studies where ectopic expression of EAAT2 in gliomas limits their growth both in vitro and in vivo. However, this concept is yet to be translated to the clinic (Robe et al. 2004).

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# **Glutamate Neurotoxicity Related** to Energy Failure

# Arne Schousboe

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#### Abstract

Glutamate is the major excitatory neurotransmitter mediating its effects via a plethora of receptors being either ionotropic or metabotropic. Each of these two mechanistically different receptor subtypes can be subdivided into distinctly different subclasses based on different pharmacological properties. Under physiological conditions the glutamatergic neurotransmission is instrumental for a large number of basic neurochemical functions such as learning and memory. Therefore, pharmacological manipulations of glutamatergic neurotransmission

A. Schousboe

Department of Drug Design and Pharmacology, Faculty of Health and Medical Sciences, University of Copenhagen, Copenhagen, Denmark e-mail: as@farma.ku.dk; arne.schousboe@sund.ku.dk

are associated with severe side effects. One particular pathophysiological area in which pharmacological intervention in glutamatergic neurotransmission has been of considerable interest is related to energy failure such as that observed during stroke leading to an ischemic condition. This has been shown to be associated with a large overflow of glutamate into the extracellular space of the brain which leads to over activation of glutamate receptors resulting in massive neuronal degeneration normally referred to as excitotoxicity. The reason for this overflow of glutamate is the fact that efficient removal of glutamate from the extrasynaptic area is mediated by a number of highly efficient, highaffinity glutamate transporters, the majority of which is located on astrocytes ensheathing the synapses. As the transporters are functionally coupled to the Na<sup>+</sup>/K<sup>+</sup>-ATPase, energy failure leading to reduced levels of ATP renders the transporters functionally inadequate resulting in efflux of glutamate from the cytoplasmic pool of glutamate. The energy substrates in the brain are under normal conditions limited to glucose and lactate but also glycogen which is selectively located in the astrocytes can play an important role both under physiological and pathophysiological conditions. These aspects are discussed in detail and evidence is presented pointing towards a hitherto neglected role of glycogen in the maintenance of glutamatergic activity during physiological conditions.

List of Abb	previations
AAT	Aspartate aminotransferase
AMPA	(RS)-2-amino-3-(3-hydroxy-5-methyl-4-isoxazolyl)propionic acid
EAAT-1	Excitatory amino acid transporter-1
EAAT-2	Excitatory amino acid transporter-2
GABA	Gamma-amino butyric acid
GDH	Glutamate dehydrogenase
GLAST	Glutamate-aspartate transporter
GLT-1	Glutamate transporter-1
GS	Glutamine synthetase
KA	Kainic acid
LDH	Lactate dehydrogenase
NMDA	N-methyl-D-aspartic acid
PAG	Phosphate-activated glutaminase
PC	Pyruvate carboxylase
TCA	Tricarboxylic acid

# 1 Introduction

Energy failure in the brain may be the result of stroke or global ischemia, both of which will lead to a drastic reduction of the availability of glucose and oxygen (Dugan and Kim-Han 2006). This condition is normally referred to as

oxygen-glucose deprivation (OGD) when used in in vitro systems such as cultured cells (e.g., Bonde et al. 2003). This condition is associated with degeneration of neurons in the brain as well as in tissue culture systems. It was shown in the pioneering work of Lucas and Newhouse (1957) that exposure of the central nervous system to excess amounts of glutamate will lead to neurodegeneration. Subsequently it was demonstrated that ischemia results in overflow of glutamate to the extracellular space in the brain (Benveniste et al. 1984). Based on this, it was suggested that glutamate would be likely to play a central role in the mechanisms leading to neurodegeneration induced by ischemia. The term excitotoxicity was coined by Olney et al. (1971) based on the observation that glutamate is not only an excitatory neurotransmitter (Curtis et al. 1959) but also acts as a neurotoxin.

The present review will deal with a delineation of the role of glutamate receptors in the neurotoxic actions of glutamate as well as a discussion of mechanisms involving glutamate transporters responsible for the overflow of glutamate during energy failure. This will include a discussion of glutamate metabolism as well as the role of glycogen, glucose, and lactate as energy substrates in relation to glutamate neurotransmission. Additionally, it will be discussed to what extent inhibitory neurotransmission mediated by GABA could play a role as a regulatory mechanism.

# 2 Glutamate Receptors

#### 2.1 Ionotropic Receptors

Based on pharmacological characterization and subsequent cloning, the ionotropic glutamate receptors are subdivided into three major classes, i.e., N-methyl-D-aspartic acid (NMDA), (*RS*)-2-amino-3-(3-hydroxy-5-methyl-4-isoxazolyl) propionic acid (AMPA), and kainic acid (KA) receptors (Braüner-Osborne et al. 2000). As each of these subgroups consists of a number of subunits that can be assembled in different combinations, the total number of possible receptors is much higher and these different receptors have pharmacologically and functionally distinct properties (Braüner-Osborne et al. 2000). This has turned out to be very important in attempts to design strategies to ameliorate glutamate-mediated neurodegeneration (Schousboe and Frandsen 1995; Ikonomidou and Turski 1995; Braüner-Osborne et al. 2000).

# 2.2 Metabotropic Glutamate Receptors

Subsequent to the demonstration that exposure of nervous tissue preparations to glutamate resulted in stimulation of inositol phospholipid metabolism (Sladeczek et al. 1985; Nicoletti et al. 1986a, b), cloning studies have identified eight subtypes of metabotropic glutamate receptors (see Braüner-Osborne et al. 2000). These are further subdivided into three subgroups consisting of group I acting on

phospholipase C leading to an increase in inositol phosphates and groups II and III that are coupled to adenylate cyclase which is inhibited leading to a decrease in the cellular content of cyclic AMP (see Braüner-Osborne et al. 2000). Due to the fact that this group of receptors plays a fundamental role in the regulation of a number of important brain functions, an enormous effort has been made to make glutamate analogs of restricted conformation acting as agonists and antagonists on the different subtypes and a comprehensive list of such compounds is provided by Bräuner-Osborne et al. (2000).

# **3 Glutamate Transporters**

Subsequent to its release as neurotransmitter and interaction with the receptors, glutamate must be inactivated. This process is mediated via high-affinity transporters, the capacity of which is more pronounced in astrocytes compared to neurons (Schousboe 1981; Gegeslashvili and Schousboe 1997; Danbolt 2001; Schousboe et al. 2012). This notion is based on studies of the kinetics of glutamate uptake into cultured neurons and astrocytes as well as detailed immunocytochemical localization studies using specific antibodies prepared from peptide libraries generated from the cloned transporters (see Schousboe et al. 2012 and references therein). The following sections will provide a short description of the two major astroglial transporters GLAST (EAAT-1) and GLT-1 (EAAT-2).

# 3.1 GLAST and GLT-1

A glutamate transporter was co-purified with the enzyme UDP galactose: ceramide galactosyltransferase (Schulte and Stoffel 1995) and this procedure was used to clone the glutamate-aspartate transporter named GLAST (Storck et al. 1992). At the same time another glutamate transporter was purified using a synaptosomal membrane preparation from rat brain (Danbolt et al. 1990). This enabled the production of a specific antibody which was subsequently used to clone this glutamate transporter named GLT-1 as it turned out to be different from GLAST (Pines et al. 1992). In spite of the fact that synaptosomal membranes had served as the starting material in the purification procedure, it became clear from immunohistochemical studies at the light and electron microscopic level that GLT-1 is almost exclusively localized in astroglial cells (Danbolt et al. 1992; Levy et al. 1993; Rothstein et al. 1994; Chaudhry et al. 1995). While GLAST is also essentially only expressed in astrocytes (Chaudhry et al. 1995; Lehre et al. 1995; Schmitt et al. 1997), the regional expression pattern of the two transporters shows a differential distribution with higher expression of GLT-1 in cortical areas and higher levels of expression of GLAST in cerebellum compared to cortex (Lehre et al. 1995).

# 3.2 Energy Requirements of Glutamate Transport

Detailed studies by Levy et al. (1998) of the stoichiometry of the ionic dependence of glutamate transport mediated by GLT-1 have shown that this requires co-transport of three sodium ions and one proton and countertransport of one potassium ion. This movement of ions needs to be reversed by the concerted action of the Na<sup>+</sup>/K<sup>+</sup>-ATPase and the Na<sup>+</sup>/H<sup>+</sup> exchanger. As the latter process is also dependent on the ATPase, a total of four sodium ions need to be transported which consumes 1.3 mol of ATP. Interestingly, it has been reported recently that a tight coupling appears to exist between the glutamate transporters and the sodiumpotassium ATPase (Cholet et al. 2002; Rose et al. 2009; Genda et al. 2011; Bauer et al. 2012) which would make the processes highly efficient.

It should be considered that the neurotransmission process consists of not only inactivation of glutamate by the astrocytic transporters but also of the vesicular packaging of glutamate prior to its release as a transmitter. This process involves the vesicular H<sup>+</sup>-ATPase (Maycox et al. 1988) which likely has the same ATP stoichiometry as the mitochondrial F-type ATPase coupled to ATP production. Hence, vesicular filling is associated with approximately one third mol of ATP per mol of glutamate transported into the vesicles.

# 4 Glutamate Metabolism

#### 4.1 Astrocytes

Glutamate transported into the astrocytes subsequent to release from vesicles in the presynaptic nerve endings will be metabolized either by the astrocyte specific enzyme glutamine synthetase (GS) or by oxidation in mitochondria after conversion to  $\alpha$ -ketoglutarate catalyzed by either glutamate dehydrogenase (GDH) or aspartate aminotransferase (AAT) as described in detail elsewhere (e.g., Sonnewald et al. 1997; Schousboe et al. 2012). There is evidence to suggest that the metabolic pathways involved in the oxidation of glutamate are dependent on the prevailing glutamate concentration in the astrocyte (Sonnewald et al. 1997). Thus, the GDH catalyzed conversion to  $\alpha$ -ketoglutarate prevails over the transamination catalyzed by AAT at elevated glutamate concentrations. This impacts on the energy homeostasis of glutamate metabolism since the GDH reaction produces NADH which subsequently gives rise to ATP production by oxidative phosphorylation (see McKenna et al. 2012), whereas the transamination reaction does not provide energy in the form of ATP. Regardless of the reaction responsible for the conversion of glutamate to  $\alpha$ -ketoglutarate, subsequent metabolism of this tricarboxylic acid (TCA) cycle intermediate to carbon dioxide via pyruvate recycling (see Waagepetersen et al. 2009) provides 24 mol of ATP. Hence, glutamate can be regarded as an excellent energy substrate in the brain and its metabolism can to some extent replace that of glucose in astrocytes (Hertz et al. 2000).

#### 4.2 Neurons

In both glutamatergic and GABAergic neurons, glutamate is formed by deamination of glutamine catalyzed by phosphate-activated glutaminase (PAG). Glutamine must be provided by surrounding astrocytes as neurons are incapable of performing a net synthesis of glutamine from glucose as they do not express the enzymes pyruvate carboxylase (PC) and GS (see Schousboe et al. 2012), both of which are astrocyte specific (Norenberg and Martinez-Hernandez 1979; Yu et al. 1983). In glutamatergic neurons this newly formed glutamate can be taken up into vesicles for transmitter use or it may be used as an energy substrate as explained above (Sect. 4.1). In GABAergic neurons it is either converted to transmitter GABA by decarboxylation or used as energy substrate (see Schousboe et al. 2012).

# 5 Consequences of Energy Failure in Glutamatergic Neurotransmission

# 5.1 Glycogen, Glucose, and Lactate as Energy Substrates

The brain is the most energy-demanding organ accounting for about 20 % of the resting metabolic rate albeit amounting to only 2 % of the body weight (see McKenna et al. 2012). The majority of the energy consumption (75 %) is related to signaling processes, whereas the remaining energy consumption is associated with basic cellular activities (McKenna et al. 2012). In the adult brain glucose is the main obligatory energy substrate but alternative substrates such as ketone bodies and lactate can be utilized. The use of ketone bodies will be prominent during prolonged fasting (McKenna et al. 2012). Interestingly, during strenuous exercise leading to lactate production in the muscles and high lactate plasma levels, lactate can be taken up into the brain (Quistorff et al. 2008). Glucose is metabolized in the glycolytic metabolic pathway leading to production of pyruvate which can either be oxidatively metabolized in the mitochondria to CO<sub>2</sub> and water or it can be reduced to lactate catalyzed by lactate dehydrogenase (LDH) in which case lactate is the end product of glucose metabolism. It has been proposed that in astrocytes glucose is preferentially converted to lactate which is subsequently transferred to the neurons where it is converted to pyruvate via the LDH reaction and further oxidatively metabolized (Pellerin and Magistretti 1994). This process, referred to as the astrocyte-neuron lactate shuttle, has subsequently been extensively discussed and it seems clear that astrocytes have a significant oxidative metabolism which on a volume basis is comparable to that in neurons (Hertz et al. 2007). It is important to note that the combination of glycolysis and oxidative metabolism of pyruvate requires the presence of the malate-aspartate shuttle (or an equivalent mechanism) which can secure exchange between the cytosol and the mitochondria of the redox couple NAD<sup>+</sup>/NADH as the mitochondrial membrane is impermeable to NAD<sup>+</sup> and NADH (McKenna et al. 2012). In this context it should be noted that astrocytes are equipped with all enzymes as well as mitochondria necessary to carry out the

metabolic processes involved in oxidative metabolism of glucose to  $CO_2$  and water (Lovatt et al. 2007). It is also interesting to note that lactate is unable to fully replace glucose as an energy substrate for neurons during enhanced activity as that seen during exposure to depolarizing signals in the form of glutamate or elevated potassium (Bak et al. 2006; 2012).

During recent years brain glycogen has attracted attention although its concentration is quite low compared to that in other organs such as liver and muscle. However, its concentration exceeds that of free glucose, but during glucose deprivation this amount of glycogen would, if metabolized by the concerted action of glycogenolysis and glycolysis, only provide enough energy to maintain brain function for a few minutes (McKenna et al. 2012). The availability of specific pharmacological tools to inhibit the key enzyme in glycogenolysis, glycogen phosphorylase, has facilitated studies aimed at elucidating the functional role of glycogen (Obel et al. 2012). Such studies have provided evidence that glycogen turnover involving the glycogen shunt, i.e., the conversion of glucose-6-phosphate to glycogen and its reappearance as glucose-1-phosphate and thus glucose-6phosphate (see Schousboe et al. 2010; Kreft et al. 2012), is of fundamental importance for the maintenance of a fully functional glutamatergic neurotransmission system (Schousboe et al. 2007, 2010; Obel et al. 2012; Parpura et al. 2012). Since the functional integrity of this system is a prerequisite to avoid excitotoxic neuronal death, glycogen metabolism is obviously also important in relation to neurotoxic damage (Obel et al. 2012).

#### 5.2 Energy Failure and Glutamatergic Neurotransmission

As pointed out above (Sect.3.2), glutamate transport is energy dependent due to the requirement of an intact transmembrane sodium gradient. In addition to this, the process responsible for packaging of transmitter glutamate into synaptic vesicles is ATP dependent since it utilizes the proton gradient generated by the action of the  $H^+$ -ATPase. It is interesting that the ATP required in both of these processes may originate from the glycolytic metabolic pathway rather than the TCA-coupled oxidative phosphorylation pathway (Ikemoto et al. 2003; Schousboe et al. 2011). As pointed out above, glycogen metabolism and turnover seems to play an important functional role in this context.

Regardless of the metabolic processes required for ATP production, it is clear that failure of generation of ATP will result in a reduced glutamatergic transmission, i.e., vesicular release will be decreased (Schousboe and Waagepetersen 2005). Since the GABA-mediated inhibitory neurotransmission is based on the same principles as the glutamatergic system, GABA neurotransmission will be reduced as well. A possible functional interrelationship regarding glutamatergic and GABAergic neurotransmission will be provided below (see Sect. 6). As pointed out above the high-affinity glutamate transporters are energy dependent and as a consequence of this, energy failure will result in reversal of the transporters and an overflow of glutamate from the metabolic, cytosolic pool into the extracellular

space (see Schousboe and Waagepetersen 2005). This, in turn, will lead to an overstimulation of glutamate receptors resulting in excitotoxic neurodegeneration (see Sect. 5.3).

#### 5.3 Energy Failure and Excitotoxicity

The demonstration almost 30 years ago by Benveniste et al. (1984) using the microdialysis technique that ischemia leads to an excessive overflow of glutamate and aspartate in rat hippocampus provided compelling evidence that ischemia-induced energy failure leading to pronounced neurodegeneration was very likely related to an excitotoxic mechanism as described by Olney and Ho (1970). Numerous studies using pharmacological tools to delineate the role of the different types of glutamate receptors (see Sect. 2) performed in cultured neurons as well as in the intact brain in vivo during the years subsequent to the study by Benveniste et al. (1984) have demonstrated that overstimulation of these receptors is responsible for the neurodegeneration associated with brain energy failure (see Choi and Rothman 1990; Schousboe and Frandsen 1995).

In accordance with the fact that glutamate transporters are energy dependent, energy failure has been shown to result in reversal of the glutamate transporters (Nicholls and Attwell 1990; Phillis et al. 2000; Bonde et al. 2003). This therefore is the most likely explanation why ischemia leads to overflow of glutamate and aspartate as mentioned above.

The molecular mechanisms associated with stimulation of glutamate receptors subsequently leading to neuronal degeneration are extremely complex involving disturbances of  $Ca^{2+}$  homeostasis, free radical formation, and NO generation, the concerted action of which results in activation of proteases, DNA degradation, and disruption of the cytoskeleton (see, e.g., Dugan and Kim-Han 2006). A detailed discussion of this is beyond the scope of the present review.

# 6 Possible Interaction Between Excitatory and Inhibitory Neurotransmission

There is evidence to suggest that the expression level of GABA<sub>A</sub> receptors may have an impact on glutamatergic, excitatory activity and excitotoxicity is exacerbated by reduced GABA receptor activity (Schwartz-Bloom and Sah 2001). Additionally, the metabotropic GABA<sub>B</sub> receptors are involved in the regulation of glutamate release and hence excitatory transmission (Nicoll 2004). In this context it is interesting that cerebral ischemia and oxygen-glucose deprivation, both of which will result in energy failure, result in a decrease in the cell surface expression of GABA<sub>A</sub> receptors (Alicke and Schwartz-Bloom 1995; Mielke and Wang 2005) which as stated above contributes to ischemic brain damage (Schwartz-Bloom and Sah 2001). This interrelationship between glutamatergic and GABAergic neurotransmission is of importance not only for neurodegeneration induced by energy failure but also for generation of seizures and epileptic episodes. A recent review provides a detailed discussion of the significance of the machinery for synthesis, release, transport, and metabolism of glutamate and GABA as targets for development of drugs for seizure control (Rowley et al. 2012). This may be interesting in relation to neurodegeneration as well.

# 7 Conclusion

The present short review has provided a discussion of the correlation between energy metabolism and the homeostatic mechanisms regulating excitatory, glutamatergic activity. It is clear that energy failure will inevitably result in excessive glutamate overflow which, in turn, will produce neurodegeneration. Three decades of intensive pharmacological research has provided a wealth of information about strategies to ameliorate such neuronal damage but little has so far been achieved regarding development of clinically efficient drugs to prevent ischemia-induced neurodegeneration.

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# Glutamatergic Receptors in Parkinson's Disease

A. Sampedro, E. Tarragón, J. E. Yuste, F. Ros-Bernal, V. Ortiz, C. M. Campuzano, A. Gómez, C. M. Ros, E. Fernández-Villalba, and M. T. Herrero

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A. Sampedro • V. Ortiz • A. Gómez • E. Fernández-Villalba

M.T. Herrero (🖂)

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain

e-mail: ertarr@um.es; jeyuste@um.es; pacoros@um.es; carmenmariaros@um.es; mtherrer@um.es; ezquerro@med.uji.es

Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain e-mail: vicortiz@um.es; aurora.gomez@um.es

E. Tarragón • J.E. Yuste • F. Ros-Bernal • C.M. Campuzano • C.M. Ros •

Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

#### Abstract

Extracellular concentrations of the excitatory glutamate are maintained at relatively low levels to avoid an excessive activation of glutamate receptors (GluRs) that can trigger a cell death. This neuronal loss induced by hyperexcitation of GluRs has been associated with a wide range of acute and chronic neurological disorders such as Parkinson's disease (PD). In fact, altered glutamatergic neurotransmission and metabolical dysfunction seems to be crucial in the pathophysiology of PD. Moreover, degeneration of dopamine nigral neurons of the substantia nigra pars compacta (SNpc) provokes striatal dopaminergic denervation and a cascade of functional modifications in the activity of basal ganglia nuclei including excitatory glutamate transmission. Glutamate receptors are divided into two distinct groups, ionotropic and metabotropic receptors. The ionotropic receptors (iGluRs) are further subdivided into three groups: (i) α-amino-3-hydroxy-5-methyl-4-isoxazolepropionate (AMPA), (ii) kainate, and (iii) N-methyl-D-aspartate (NMDA) receptor channels. The metabotropic receptors (mGluRs) are coupled to GTP-binding proteins (G proteins) and regulate the production of intracellular messengers. In this chapter, we review recent progress in the research of GluRs with special emphasis on the molecular diversity of the mGluR system and its implications in the physiopathology of PD. Finally, based on these evidences, we highlight possible therapeutic strategies that might be important to slow down the progression of PD as well as to modulate levodopa-induced dyskinesias.

#### Keywords

AMPA • GluRs • Glutamate • Neurotoxicity • NMDA • Parkinson's disease • Substantia Nigra

#### 1 Introduction

Parkinson's disease (PD) is a neurodegenerative disorder characterized by motor symptoms including tremor and bradykinesia (Dauer and Przedborski 2003; Schapira 2009). The primary pathophysiology underlying PD is the degeneration of dopaminergic neurons in the substantia nigra *pars compacta* (SNpc) (Jellinger 2002). Loss of these neurons causes pathological changes in neurotransmission in the basal ganglia circuits (Carlsson et al. 1957). The ability of ionotropic and metabotropic glutamate receptors to modulate neurotransmission throughout the basal ganglia suggests that these receptors may be targets for reversing the effects of altered neurotransmission in PD (Blandini and Armentero 2012; Leaver et al. 2008). Studies in animal models suggest that modulating the activity of these receptors may alleviate the primary motor symptoms of PD as well as the side effects induced by dopamine replacement therapy (Gardoni et al. 2012; Konitsiotis et al. 2000). Moreover, glutamate receptor ligands may slow disease progression by delaying progressive dopamine neuron degeneration

(Fell et al. 2009; Johnson et al. 2009). Antagonists of N-methyl-D-aspartate (NMDA) receptors have shown promise in reversing motor symptoms, levodopainduced dyskinesias, and neurodegeneration in preclinical PD models (Blandini and Armentero 2012; Johnson et al. 2009). The effects of drugs targeting  $\alpha$ -amino-3hydroxy-5-methyl-4-isoazolepropionic acid (AMPA) receptors are more complex; while antagonists of these receptors exhibit utility in the treatment of levodopainduced dyskinesias, AMPA receptor potentiates show promise for neuroprotection (Johnson et al. 2009; Kobylecki et al. 2010). Pharmacological modulation of metabotropic glutamate receptors (mGluRs) may hold even more promise for PD treatment due to the ability of mGluRs to fine-tune neurotransmission. Antagonists of mGluR5, as well as activators of group II mGluRs and mGluR4, have shown promise in several animal models of PD (Johnson et al. 2009). As these drugs reverse motor deficits in addition to providing protection against neurodegeneration, glutamate receptors therefore represent exciting targets for the development of novel pharmacological therapies for PD. However, several approaches have been undertaken concluding in few results remarkable for humans. Nonselective blockade induces unacceptable psychiatric and cognitive adverse effects, while selective blockade is still under investigation with little benefit to date (Fantin et al. 2008; Hallett and Standaert 2004). The arduousness of finding an appropriate pharmacological therapy targeting NMDA receptors prompted much interest on metabotropic glutamate receptors (mGluRs) as novel targets for the management of PD.

# 2 Glutamate Receptor Subtypes

#### 2.1 Ionotropic Glutamate Receptors

There are three major types of ionotropic glutamate receptors (iGluRs), which are named after the agonists that were originally identified to activate them selectively, and are, thus, called NMDA, AMPA, and 2-carboxy-3-carboxymethyl-4isopropenylpyrrolidine (kainate) receptors (Fonnum 1984; Kostandy 2012). Native receptors of all of these families are likely heteromeric assemblies comprising more than one type of subunit (Kew and Kemp 2005). Although the average overall amino acid identity of ionotropic glutamate receptor subunits across the three families is only in the 20–30 % range, they share common structural features, which clearly place them into a single large superfamily. The transmembrane topology of the glutamate receptor family differs from the classic four transmembrane domain model of the nicotinic acetylcholine receptor, which has extracellular amino- and carboxy-termini and a large intracellular loop between transmembrane domains 3 and 4 (Traynelis et al. 2010). Ionotropic glutamate receptor subunits possess an extracellular amino-terminal domain, which exhibits homology to the metabotropic glutamate receptor (mGluR) bilobed agonist-binding domain, followed by a first transmembrane domain, and then a pore-forming membraneresiding domain that does not cross the membrane but forms a reentrant loop entering from and exiting to the cytoplasm (Fonnum 1984). The second and third transmembrane domains are linked by a large extracellular loop, and the third transmembrane domain is followed by an intracellular carboxy-terminus (Dingledine et al. 1999; Mayer and Armstrong 2004). The crystal structure of the iGluR ligand-binding domains, which comprise polypeptides in both the amino-terminus (S1 domain) and the extracellular loop between transmembrane domains 3 and 4 (S2 domain), has confirmed this topology model for the ionotropic gluta-mate receptor family (Armstrong et al. 1998; Mayer and Armstrong 2004). Several studies have shown an implication of NMDA and AMPA receptors in PD alterations but not kainate receptors, so we haven't focused on them in this chapter.

#### 2.1.1 NMDA Receptors

*N*-methyl-D-aspartate (NMDA) glutamate receptors are excitatory receptors widely involved in the basal ganglia circuitry and therefore constitute a promising target for the development of new drugs to prevent or treat PD (Hallett and Standaert 2004). NMDA receptors are ligand-gated ion channels composed of multiple subunits, which have distinct cellular and regional patterns of expression. NMDA receptors are widely found in all of the structures implicated in basal ganglia circuits, although there is a different composition of its units depending on the physiology of the circuit. For example, NMDA receptors present in the striatum, which mainly are located postsynaptically on striatal neurons, are crucial for dopamine-glutamate interactions since the abundance, structure, and function of these striatal receptors are modified by dopamine depletion and they could be further varied by pharmacological treatments used in PD (Hallett et al. 2005). In animal models, NMDA receptor antagonists are effective antiparkinsonian agents and can reduce the complications of chronic dopaminergic therapy (wearing off and dyskinesias) (Del Dotto et al. 2001). Use of these agents in humans has been limited because of the adverse effects associated with nonselective blockade of NMDA receptor function, but the development of more potent and selective pharmaceuticals holds the promise of important new therapeutic approach for PD.

#### 2.1.2 AMPA Receptors

AMPA receptors are mainly implicated in most of the excitatory neurotransmission in the brains of mammals and also taking part in synaptic plasticity processes thought to form the base of memory and learning and the organization of neural networks during embryogenesis (O'Neill and Witkin 2007). It has been shown, using molecular cloning techniques, that the AMPA receptor family has a group of four different subunits named GluR1–4 being native AMPA receptors most likely tetramers produced by the assembly of one or more of these subunits generating homomeric or heteromeric receptors (Santos et al. 2009). Experimental data indicate that the modulation of AMPA receptors, correlated with an increased hippocampal activity, could have a crucial role in the therapeutic treatment of cognitive deficits (Chappell et al. 2007). Moreover, clinical studies have indicated that AMPA receptor modulators enhance cognitive function not only in elderly subjects but even in neurological and psychiatric patients (Lees 2000). The pathway in which AMPA receptors is implicated

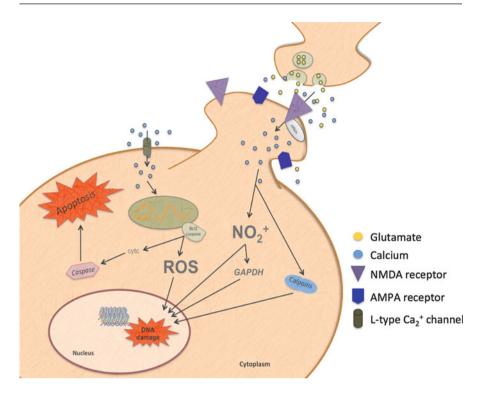
is still known, but it has been postulated, using in vitro and in vivo studies, that AMPA receptors could induce the activation of a protein tyrosine kinase, Lyn, which would recruit the mitogen-activated protein kinase (MAPK) signaling pathway increasing the expression of BDNF (Hayashi et al. 1999; Zhang et al. 2010). In conclusion, several selective and systemically active AMPA receptor potentiators have been reported which main functions are to modulate glutamatergic transmission, to enhance synaptic transmission and long-term potentiation, and to increase neurotrophin expression (O'Neill et al. 2004). All these functions situate AMPA receptor potentiators and depression and stimulating neuroprotective effects in PD.

### 2.2 Metabotropic Receptors

Glutamate receptors are a group of G protein-coupled receptors (GPCRs), which are differentially localized throughout the basal ganglia and could be a good therapeutic target for PD treatment (Waxman et al. 2007). Glutamate receptors regulate excitatory and inhibitory synaptic transmission acting on synaptic mechanisms (Waxman and Lynch 2005). Structurally, we can difference eight subtypes of mGluRs, divided into three groups according to their sequence homology and G protein-coupling specificity (Conn et al. 2005; Conn and Pin 1997; Pin and Acher 2002; Pin and Duvoisin 1995). Group I mGluRs include mGluR1 and mGlu5, both of them activate phospholipase and protein kinase C and raise intracellular calcium levels. Group II mGluRs consist of two mGluR subtypes (GluR2 and GluR3) which mainly inhibit adenylyl cyclase. Group III mGluRs include mGluR4, mGluR6, mGluR7, and mGluR8, which structurally are located presynaptically, and functionally act causing a reduction in neurotransmitter release (Waxman et al. 2007). mGluRs are the most relevant glutamate receptors implicated in therapeutic strategies for PD treatment at present and will be widely reviewed in this work.

# 3 Distribution of Glutamatergic Receptors and Excitotoxicity

In spite of its ubiquitous role as an excitatory neurotransmitter in the intermediary metabolism in the brain, glutamate can be highly toxic to neurons, a phenomenon known as "excitotoxicity." This neuronal death is a result of an excessive stimulation of the NMDA receptor and the subsequent massive influx of extracellular  $Ca^{+2}$ . This increase in cytoplasmic  $Ca^{+2}$  activates a number of  $Ca^{+2}$ -dependent enzymes implicated in the catabolism of phospholipids, proteins, nucleic acids, and the nitric oxide synthase activity. All this events lead to a global breakdown of cellular homeostasis including membrane rupture, cytoskeletal alterations, and nitric and oxide free-radical production (Koutsilieri and Riederer 2007) (Fig. 1). During the last decade numerous reviews have addressed the role of glutamate receptors and related synaptic mechanisms in preclinical research on neurodegenerative diseases



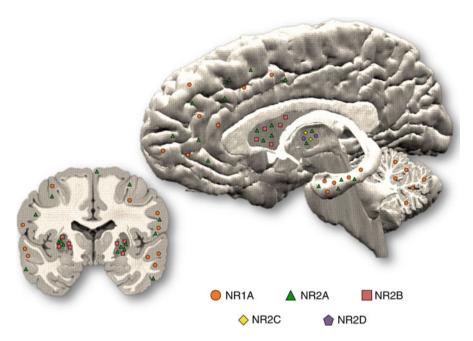
**Fig. 1** Excitotoxicity of glutamate release and its implication in synaptic plasticity. One of the major functions of glutamate receptors, both metabotropic and ionotropic receptors, appears to be the modulation of synaptic plasticity. Overstimulation of NMDA receptor increases  $Ca^{2+}$  influx into the cell (neuron) which leads to the breakdown of cellular homeostasis, the lost of mitochondria function, and the release of nitric oxide and other free radicals

(Johnson et al. 2009; Lau and Tymianski 2010; Luscher and Huber 2010; Ondrejcak et al. 2010; Palop and Mucke 2010a, b; Randall et al. 2010).

# 3.1 Ionotropic Receptors

#### 3.1.1 NMDA Receptors

Studies designed to determine NMDA receptor distribution have shown that these receptors are found throughout the brain but predominantly within the forebrain (Bean et al. 2006; Beckerman et al. 2012; Collingridge et al. 2013; Monaghan et al. 1998) (Fig. 2). Furthermore, quantitative comparisons of the distribution of NMDA receptors determined by the binding sites of the different ligands have suggested the presence of multiple pharmacologically distinct types of NMDA receptors (Monaghan and Jane 2009). Following the molecular cloning of NMDA receptor subunits, the distribution of each subunit was examined using in situ hybridization, histochemistry, and biochemical analysis using subunit-specific antibodies



**Fig. 2** Distribution of ionotropic glutamate receptors in the human brain. Using the binding sites of different ligands, the ubiquitous distribution of ionotropic glutamate receptors in the nervous system especially in the forebrain has been determined

(Dunah et al. 1999; Yamada and Nabeshima 1998). The NR1A mRNA is distributed widely in the brain but more so in the cerebral cortex, hippocampus, and cerebellum (Ehlers et al. 1995; Hara and Pickel 2008). The NR2A transcript is selectively present in the forebrain with a high-level expression in the cerebral cortex, hippocampus, septum, caudate-putamen, and olfactory bulb (Ling et al. 2012). The NR2B mRNA is expressed predominantly in the granule cell layer of the cerebellum, with weak expression in the olfactory bulb and thalamus (Jarzylo and Man 2012; Kaufmann et al. 2012). In sum, while NR2A and NR2B transcripts are found in a subset of hippocampal neurons, most likely interneurons (Kuppenbender et al. 2000), low levels of the NR2D transcript are detected in the thalamus, brain stem, and olfactory bulb, being predominant in the supraoptic nucleus (Doherty and Sladek 2011).

The location of NMDARs is also related with its signaling: (i) intrasynaptic NMDAR location is involved in neuronal survival, and (ii) extrasynaptic NMDARs have a crucial role in cell-death pathways (Papadia et al. 2005). Among other neurological alterations, NMDA receptor dysfunction may be a primary factor in the aetiology of schizophrenia, where NMDA receptor antagonists induce cortico-limbic degeneration and induct psychotic states, thus hastening a neurotoxicity and neuronal degeneration, which can be hampered using dopaminergic antagonists (Dawson et al. 2012).

# 3.1.2 AMPA Receptors

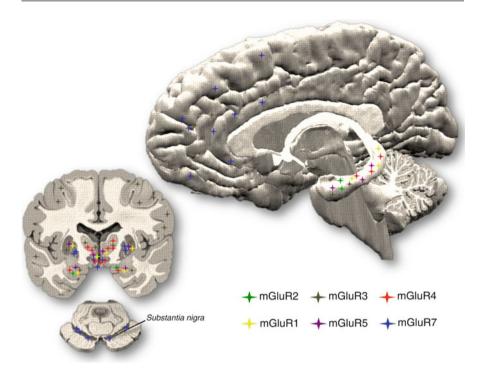
It has been shown that AMPA receptors in glutamatergic synapses mediate the transmission of low frequency and are involved in the expression of long-term potentiation (LTP) and long-term depression (LTD) processes, considered the cellular correlates of memory formation (Bliss and Collingridge 1993). At early time points of human embryogenesis, most of excitatory synapses are thought to be postsynaptically "silent," since their functional receptors are NMDA receptors but not AMPA ones (Daw et al. 2007; Isaac et al. 1995; Li and Zhuo 1998; Liao et al. 1995; Scott et al. 2007). Such selective pattern of synaptic activation has been termed "silent synapses" because the NMDA pore is largely hampered at resting membrane potentials (Isaac et al. 1995; Nowak et al. 1984; Scott et al. 2007). Following protocols that generate LTP, however, these functionally inhibited synapses rapidly become AMPA receptor responsive (Isaac et al. 1995; Liao et al. 1995). One possible mechanism to explain this observation is a synaptic transfer of a pool of preassembled AMPA receptors from an intracellular compartment to the postsynaptic membrane. Then, this intimate relation between AMPA and NMDA receptor is the basis of the search of alternative pathways to prevent the activation derived from an excessive NMDA excitotoxicity: NMDA antagonists, as AMPA-type receptors, have been tested and have demonstrated beneficial effects on the damage to the white substance covering the axons of neurons (Park et al. 2003). Its efficacy in preclinical models of stroke is greater than some NMDA receptors' antagonists, but some of these compounds exhibit problems of solubility or adverse effects, primarily by inhibiting the fast excitatory synaptic transmission (Clarkson et al. 2011). Excitotoxicity mediated by the alterations of AMPA receptor has been implicated in the selective vulnerability of motor neurons in amyotrophic lateral sclerosis (ALS) (Corona and Tapia 2007), in the induction of neurological deficits including schizophrenia (Tsai and Coyle 2002), and it could also underlie the cognitive deficits related with neurodegenerative diseases such as Alzheimer's disease (Kamenetz et al. 2003).

# 3.2 Metabotropic Receptors

Metabotropic receptors (mGluRs) are expressed throughout the CNS (Fig. 3). However, specific subtypes are expressed in restricted areas, such as the hippocampus where preferential expression of the receptor subtypes mGluR1<sub>b</sub>, mGluR2/3, mGluR4<sub>a</sub>, and mGluR5 has been demonstrated (Blumcke et al. 1996). mGluR3 is expressed widely in the CNS (Makoff et al. 1996a, b). mGluR4 expresses different subtypes in the thalamus, hypothalamus, and caudate nucleus, and mGluR6 is expressed in the retina (Vardi et al. 2000) (Fig. 2).

# 3.2.1 Group I mGluRs

Both mGlu1 and mGlu5 are most often located postsynaptically, where they are related to neuronal depolarization and excitability (Lujan et al. 1997). Nevertheless, in the midbrain, mGluR1 induces hyperpolarization of dopamine neurons



**Fig. 3** Distribution of metabotropic glutamate receptors in the human brain. Metabotropic receptors are ubiquitously localized throughout the CNS, although its different subtypes are specifically expressed in the hippocampus, thalamus, and striatum

(Katayama et al. 2003). Both receptors are primarily coupled to Gq/G11 proteins (Masu et al. 1991) and have been implicated in some human disorders such as inherited ataxia and cerebellar disorders, associated with an altered expression of mGlu1 receptors in Purkinje cells (Catania et al. 2001; Conti et al. 2006; Sachs et al. 2007). However, although these data suggest that the mGluR1 could be implicated in inherited cerebellar disorders, no inductive alterations on this receptor have been determined in infant idiopathic early-onset ataxia patients (Rossi et al. 2010). Moreover, it has also been shown that mGlu1 receptor antagonists can protect hippocampal neurons against "postischemic" degeneration induced by oxygen-glucose deprivation (Pellegrini-Giampietro 2003). Recent studies have also proposed the blockade of mGlu1 receptors as a novel treatment for schizophrenia, but it has to be corroborated in the next years (Nicoletti et al. 2011).

On the other hand it has been postulated that mGlu5 has a synergic role with NMDA receptors, both acting in the control of the neuronal activity of prefrontal cortex and also being implicated in cognitive dysfunction associated with schizo-phrenia, being a potential treatment for schizophrenic syndromes (Conn et al. 2009).

#### 3.2.2 Group II mGluRs

Group II mGluRs, mGlu2/3 receptors, are principally present in astrocytes surrounding the neuronal soma and synapses. They also have a presynaptic distribution inhibiting neurotransmitter release and leading to neuronal resting (Liu et al. 1998). These receptors are presynaptically localized on subthalamic nucleus terminals, and their activation inhibits excitatory transmission at subthalamic synapses resulting in a reduced excitatory drive through the indirect pathway (Murray et al. 2002).

Although both receptors, mGlu2 and mGluR3, are mostly expressed in the CNS, they have differences in their location, so while mGluR2 is uniquely in the preterminal region of the axons, far from the active zone of the neurotransmitter release (Lujan et al. 1997; Ohishi et al. 1993a; Tamaru et al. 2001), mGlu3 receptor is expressed not only in presynaptic and postsynaptic sites but also in glial cells (Ohishi et al. 1993b; Tamaru et al. 2001). Functionally, both receptors are implicated in the same human disorders. Both dysfunctions in the mGlu2/3 receptors are correlated with the following: (i) anxiety, both receptors regulate the synaptic transmission and plasticity in the amygdala (Lin et al. 2005; Wang and Gean 1999); (ii) schizophrenia, the hyperglutamatergic theory of schizophrenia has been corroborated using glutamate agonist in experimental animal models, and mGluR2/3 receptor agonists have been developed as antipsychotic drugs (Aghajanian and Marek 1999); and (iii) depression, the systemic treatment with a selective mGlu2/3 receptor antagonist has beneficial effects on this mental disorder (Fevissa et al. 2010). Apart from these effects described, mGluR2/3 receptors have a special significance in neurodegenerative diseases, as the inhibition of mGlu2 receptors decreases excitotoxic death, and mGlu3 receptors are also implicated in neuroprotective pathways (Corti et al. 2007).

#### 3.2.3 Group III mGluRs

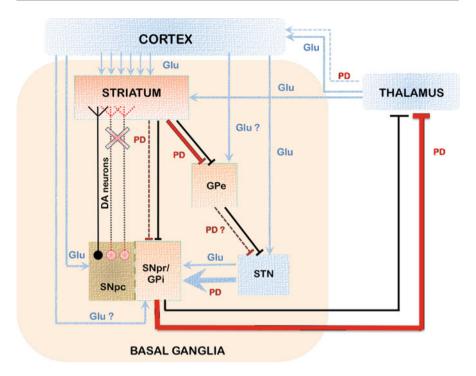
Group III mGluRs are identified in microglia and astrocytes as well as in presynaptic membrane like mGluR7 receptor (Geurts et al. 2005; Niswender and Conn 2010). This specific subcellular distribution is not still but modified during the development of the CNS and may play a key role for its proper progressive organization: (i) mGluR7 and mGluR8 are presynaptically located in the striatum, the globus pallidus, and in the substantia nigra pars reticulata (SNpr), where their activation inhibits GABA as well as glutamate transmission (Kosinski et al. 1999), and (ii) mGluR4 appears to be more selectively located in striatopallidal synapses and inhibits synaptic activity at this level (Valenti et al. 2003). In relation with this variety on location, glutamate also presents a changeable affinity for these receptors since it shows a high affinity to mGluR4 and mGluR8 but a very low one to mGluR7 (Schoepp et al. 1999). Apart from their implication in Parkinson's disease which would be further intensely described, these receptors have shown a significant relevance in human disorders such as (i) anxiety: it has demonstrated the negative modulation of both mGlu8 and mGlu4 receptors in the induced anxiety in rodents (Nicoletti et al. 2011), but mGlu7 has an uncertain role in this disease (Lavreysen and Dautzenberg 2008); and (ii) neuroinflammation and immune misbalanced: mGlu4 receptor has been recently implicated in the crosstalk between the nervous and immune system, where the activation of this receptor induces the differentiation of antigen specific T cells into T regulatory ones (Fallarino et al. 2010). With regard to mGlu6 expression is not as much ubiquitous in the CNS as the other group III receptors, and physiologically it is almost exclusively implicated in the visual region since it is localized in the dendrites of bipolar cells of the retina acting on the visual physiology in darkness and being part of human visual pathologies (Nawy 2000; Xu et al. 2012).

## 4 Implication of Glutamate Receptors in Parkinson's Disease

#### 4.1 NMDA Receptors as a Therapy for Parkinson's Disease

Under physiological conditions there is a neurotransmitter balance in striatum between the activation of striatal neurons through NMDA receptors and inhibition by the D2 receptors (Di Chiara et al. 1994). In PD, the depletion of nigrostriatal dopamine results in disinhibition of the striatal neurons and therefore in a relative glutamatergic overactivity (Greenamyre 1993). Moreover, the dopaminergic deficit results in enhanced activity of the STN which projects to the output nuclei of the basal ganglia rendering them overactive (Rodriguez et al. 1998). These known regulations classify PD as "a secondary glutamate overactivity syndrome" (Riederer et al. 2001) (Fig. 4). Changes in abundance of NMDA subunits in striatum have been described in dopamine-denervated rats as well as in MPTPtreated parkinsonian monkeys (Betarbet et al. 2004; Hallett et al. 2005). In the 6-OHDA-lesioned rat model, dopamine depletion results in alterations in abundance of NR1 and NR2B subunits in striatal synaptosomal membrane fractions, while NR2A is unchanged (Hallett and Standaert 2004). This was shown to be due to a selective reduction of complexes composed of NR1/NR2B without any change in receptors composed of NR1/NR2A. Chronic levodopa treatment of 6-OHDAlesioned rats restores the abundance of NR1, NR2A, and NR2B subunits in homogenate and synaptosomal membrane fractions (Hallett and Standaert 2004). Similar redistribution occurs in the MPTP-lesioned macaque where the dopamine depletion induced by MPTP intoxications resulted in a marked decrease in striatal NR2B subunits, while the upregulation of NR2A expression in synaptosomal membranes may play a pivotal role in levodopa-induced dyskinesias (Hallett et al. 2005).

A large number of studies have examined the efficacy of nonselective NMDA antagonists in animal models of PD showing that NMDA receptor blockade attenuates parkinsonian findings and increases dopaminergic therapy (Lange et al. 1997; Marin et al. 2000). For humans, amantadine has been the mostly studied drug in clinical trials, showing antidyskinetic properties in PD patients with motor complications as well as an increased lifespan (Hubsher et al. 2012). A related compound, memantine, which is approved for treatment of dementia, did not seem to share the efficacy of amantadine in PD (Merello et al. 1999; Varanese et al. 2010). Finally, dextromethorphan, an antitussive, reduces dyskinesias in PD patients, but patients show somnolence (Chen 2011; Jimenez et al. 1999).



**Fig. 4** Glutamatergic pathways involved in Parkinson's disease. In Parkinson's disease the striatal dopaminergic deficit results in an enhanced activity of the subthalamic nucleus which projects to the output nuclei of the basal ganglia rendering them overactive

Nowadays, based on the experimental evidence on the significance of the NMDA receptor subtypes and the limitations in the beneficial effects of the current therapy, future studies should be focused on selective NMDA antagonists in order to achieve powerful antiparkinsonian effects without adverse effects in cognition. This would facilitate cell preservation during excitotoxic processes while not causing complete inhibition of the receptor and therefore allowing physiological neurotransmission.

## 4.2 AMPA Receptors as a Therapy for Parkinson's Disease

It is difficult to define an antiparkinsonian treatment that targets AMPA receptors since they have dual effects, while AMPAR antagonists provide symptomatic benefits, AMPAR potentiators induces neuroprotection (Johnson et al. 2009). The role of AMPA receptors in dyskinesias has been well described: using animal models such as MPTP-lesioned nonhuman primates and 6-OHDA-lesioned rats, it has been demonstrated that pharmacological blockade of AMPARs decreases levodopa-induced dyskinesias (LIDs) and possibly enhances the antiparkinsonian

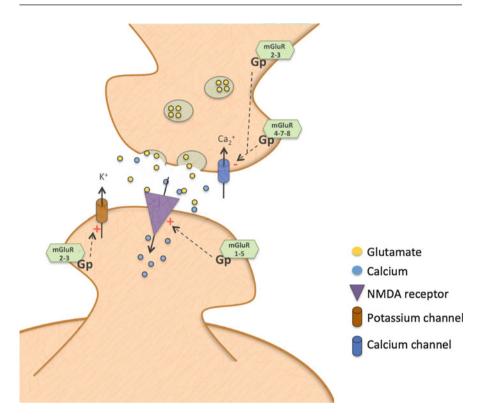
effect of levodopa (Bibbiani et al. 2005; Kobylecki et al. 2010; Konitsiotis et al. 2000). On the other hand, AMPAR agonists administered either alone or in combination with levodopa can induce dyskinesias (Konitsiotis et al. 2000). Moreover, using animal models it has been shown that the blockade of AMPA receptors could improve LIDs (Kobylecki et al. 2011; Silverdale et al. 2005). The antidyskinetic effect induced by AMPA receptor blockade derives of its action at the cellular level, as it has been demonstrated that AMPAR antagonists reduce the striatal upregulation of met-enkephalin gene expression induced by DA depletion and lately exacerbated by chronic levodopa administration (Perier et al. 2002). All these data suggest that the pathophysiological adaptations of medium spiny neurons caused by DA depletion and chronic levodopa treatment are mediated by the stimulation of AMPA receptors (Sgambato-Faure and Cenci 2012).

## 4.3 mGluRs as a Therapy for Parkinson's Disease

Several studies have shown that the pharmacological modulation of mGluRs can ameliorate motor abnormalities in experimental models of PD. There is also evidence of a neuroprotective effect against glutamatergic toxicity, to date in a lesser degree as with iGluRs. To avoid an excessive extension of the text and facilitate its comprehension, we will exclude the full-extension names of the molecules used in the different studies.

#### 4.3.1 Group I mGluRs Studies

Inhibition of group I mGluRs (mGluR1 and mGluR5) seems to be promising in PD therapy (Fig. 5). The role of group I mGluRs leading to excitotoxicity and their antagonist protective properties have been proved in non-PD models of neurodegeneration (Rao et al. 2000). There is high experimental evidence of the contribution of group I mGluRs to posttraumatic or postischemic neuronal excitotoxicity. Selective mGluR5 antagonists such as MPEP or the more selective MTEP are known neuroprotective agents. Whether this effect is mediated by mGluR5 or by off-target inhibition of postsynaptic NMDAR is controversial (Lea et al. 2005). In a striatal model of excitotoxicity, MPEP did show a neuroprotective effect, but the direct blockade of NMDA receptors appeared not to be involved (Popoli et al. 2004). The role of mGluR1 ligands in excitotoxicity is not still clear. Whereas the antagonists have shown neuroprotective activity (Faden et al. 2001), the agonists may facilitate either neurotoxicity or neuroprotection (Allen et al. 2000; Nicoletti et al. 1999). EMQMCM, a novel, highly selective uncompetitive antagonist of mGlu1 receptors, has demonstrated significant effects against kainateinduced excitotoxicity in both in vitro (in the cortical and hippocampal neuron) and in vivo animal models (Smialowska et al. 2012). In the particular case of basal ganglia, prevalent distribution of mGluR5 in the striatum and in the limbic system is well documented: both subtypes of group I mGluR are postsynaptically expressed by medium spiny neurons (Tallaksen-Greene et al. 1998) as well as by all interneurons, including cholinergic cells. As mGluRs do not cause any effect on the



**Fig. 5** Pre- and postsynaptic location of different metabotropic glutamate receptors. Among the glutamate receptors, metabotropic glutamate receptors are mostly implicated in Parkinson's disease. mGluR2 and mGluR3 are both pre- and postsynaptically located; however, mGluR4, mGluR7, and mGluR8 are located at the presynaptic level and mGluR1–mGluR5 in the postsynaptic neuron

membrane properties of the spiny neurons, its action should be indirect. Instead, they enhance the cationic currents gated by NMDA receptors (Pisani et al. 2001). Postsynaptic location of mGluR1<sub>a</sub> and mGluR5 in the subthalamic nucleus has been demonstrated showing an excitatory effect (Awad et al. 2000). PET scan of the striatum of MPTP parkinsonian monkeys showed an increased binding of mGluR5, and this upregulation of mGluR5 might play a role in disease progression in PD (Sanchez-Pernaute et al. 2008). More specifically, the blockade of mGluR5 was shown to alleviate akinesia in both 6-OHDA- and haloperidol-treated animals (Breysse et al. 2002; Ossowska et al. 2001). In addition, a study showed that mGluR5 knockout mice are less sensitive to MPTP toxicity, and mGluR5 antagonists are effective in reducing the MPTP-induced nigrostriatal damage in wild-type mice (Battaglia et al. 2003). Additionally, chronic and systemic treatment with MPEP, a selective mGluR5 antagonist, has not only anxiolytic-like effects but also neuroprotection as diminished dopaminergic cell loss in the SNpc in 6-OHDA-treated mice and rats (Chen et al. 2011).

#### 4.3.2 Group II mGluRs Studies

Group II and III mGluRs modulate the corticostriatal input which is the major excitatory drive to the basal ganglia (Fig. 5). Activation of presynaptic group II and III mGluRs inhibits glutamatergic transmission in the striatum (Rouse et al. 2000). Moreover, all three groups of mGluRs have shown to depress transmission at glutamatergic synapses in the SNpc modulating the dopaminergic pathway from the SNpc to the striatum (Wigmore and Lacey 1998). Activation of presynaptic mGluRs in the nigrostriatal synapse increases dopamine release in the direct pathway (Rouse et al. 2000). In fact, intranigral or intraventricular injection of the group II mGluRs agonist DCG-IV reduces akinesia in the reserpine-treated rat model of PD. The selective group II mGluRs agonist LY379268 was assessed to provide both functional relief and neuroprotection in the same rodent model of PD (Dawson et al. 2000). In MPTP-lesioned mice, prior treatment with LY379268 reduced the extent of nigrostriatal degeneration. Since now on, electrophysiological studies suggest that group II mGluRs are presynaptically located on STN terminals in the SNpr and the activation of these receptors selectively reduces transmission at excitatory STN synapses. Therefore, LY354740, a highly selective agonist of group II mGluRs, inhibits synaptic excitation of the SNpr (Bradley et al. 2000). However, these results were dependent on the dose and regimen of MPTP administration: (i) they were only observed under conditions in which MPTP produced a massive degeneration of nigral dopaminergic neurons, and there was no effect with lower doses of MPTP; (ii) while systemic administration of LY379268 failed to reverse akinesia or to produce typical antiparkinsonian rotational behavior, intraventricular administration provided a dose-dependent increase in locomotor activity in these rats. Moreover, the group II mGluRs antagonist, LY341495, blocked the protective effect of the LY379268 and amplified the damage produced by MPTP, even with low exposure (Battaglia et al. 2003).

#### 4.3.3 Group III mGluRs Studies

The largest evidence of clinical and pathological antiparkinsonian properties belongs to group III mGluRs. Research with general group III mGluR agonists has been developed. In this instance, intracerebral treatment with the L-SOP and L-AP4 (group III mGluR agonists) significantly reduced the akinesia induced by reserpine in rats (MacInnes et al. 2004). Injections were performed in the SNpr, GP, and intraventricularly. But the main studies have focused on specific group III mGluR subtypes.

Group III mGluRs, mGlu4R, and mGlu7R are expressed in synaptic terminals of the striatum, while there is no evidence about the presence of mGluR8. The two group III mGluR agonists ACPT-I and L-AP4 have shown similar potencies on mGluR4 and mGluR8 and lower affinity for mGluR7. No consistent effects on other mGluRs have been demonstrated. Pallidal injections of each glutamate receptor agonist have antiparkinsonian actions in haloperidol- and 6-OHDA-treated rats (Lopez et al. 2007), but the infusion of the same molecules into the SNpr had opposite effects in controls and 6-OHDA-treated rats, inducing catalepsy. Nevertheless, this akinetic effect is probably mediated by mGluR8 activation and not by mGluR4. More recently, a new group III orthosteric agonist has been identified: LSP1-3081 activates selectively and dose dependently all group III receptors transiently expressed in HEK293 cells (its potency is equivalent to that of L-AP4). LSP1-3081 was found to display the highest potency on mGlu4 receptor, followed by mGlu8, and very low on mGlu7. It is also devoid of nonspecific agonist or antagonist activity on group I or II mGluRs, and on targets other than mGluRs, it has less nonspecific effect than L-AP4 (Cuomo et al. 2009). The same authors proved that LSP1-3081 and L-AP4 depressed corticostriatal glutamatergic transmission by modulating presynaptic group III mGluRs. Both compounds also inhibited GABA input to striatal MSNs. Presumably, a reduction of glutamatergic activity on subthalamo-pallidal synapses and GABA transmission in the striatopallidal pathway are also implicated in antiparkinsonian and antiexcitotoxicity properties of group III mGluRs activation. In this sense, presynaptic activation of group III mGluRs has been shown to inhibit GABA release in the GP (Matsui and Kita 2003; Valenti et al. 2003), SNpc (Valenti et al. 2005), and SNpr (Wittmann et al. 2001). The effect of L-AP4 on striatopallidal GABAergic transmission was absent in mGluR4 KO mice (Valenti et al. 2003). These data indicate that activation of mGluR4 could decrease (or modulate) the increased inhibition of the GP that has been suggested in PD models. In the same study, direct ventricular injections of L-AP4 produced robust antiparkinsonian effect in both acute (haloperidol-induced catalepsy and reserpine-induced akinesia) and chronic 6-OHDA PD models. Taking into account the low affinity for mGluR7 of the above-described group III mGluR agonists and the fact that mGluR8 has a low distribution in the striatum, mGluR4 is probably the principal vehicle to start the response to group III mGluRs agonists. To confirm the previous hypothesis, the selective mGluR8 agonist (S)-DCPG was tested to inhibit glutamatergic signaling (Cuomo et al. 2009). However, no effects were found. The same tests were performed with L-AP4 and LSP1-3081 in mGluR4 KO mice. There was no modification of postsynaptic currents in the slices of these mice. Meanwhile, in the wild type, LSP1-3081 significantly inhibited the amplitude of postsynaptic currents. Finally, the same researchers demonstrated that intrastriatal L-AP4 injection improves akinesia in 6-OHDA-lesioned rats. The former studies emphasize the importance of subtypeselective group III mGluR agonists as antiparkinsonian drugs. However, the high conservation of the glutamate-binding site makes it difficult to develop highly selective orthosteric ligands for individual mGluR subtypes. To improve specificity, ligands that interact at sites other than the orthosteric one have been developed. These are called positive allosteric modulators (PAMs). In mice histological studies, previous systemic injection of the selective PAM of mGluR4, PHCCC, protects nigrostriatal neurons against MPTP toxicity. In knockout mice for mGlu4R, there was no benefit from PHCCC. Direct GP infusion showed the same results as systemic treatment (Battaglia et al. 2006). Unfortunately, PHCCC has low potency and poor aqueous solubility, and it is also an mGluR1 antagonist with similar potency as mGluR4 (Annoura et al. 1999). Thus, it is not useful for PD.

Using high-throughput screening (HTS), multiple novel PAMs of mGluR4 were identified. The lead from this cluster of structurally related compounds was

VU0155041. Among its particularities, it exhibits submicromolar potency; it is highly selective for mGluR4 relative to other mGluRs and other targets including NMDAR and has an improved aqueous solubility. The same researches demonstrated efficacy of VU0155041 in two rodent models of PD: haloperidol-induced catalepsy and reserpine-induced akinesia (Niswender et al. 2008). A few additional novel chemical mGluR4 PAMs have been introduced later on and have shown some antiparkinsonian activity in both haloperidol-induced catalepsy and reserpine-induced scatalepsy and reserpine-induced akinesia (Hopkins et al. 2009). One of them, VU0364770, has exhibited efficacy in reversing haloperidol-induced catalepsy, forelimb asymmetry, and attention deficits in 6-OHDA-lesioned rats. When coadministered with an adenosine 2A receptor antagonist or with levodopa, it also showed potentiation of their efficacy (Jones et al. 2012). The latest could mean a levodopa-sparing role of the compound. In conclusion, mGluR4 positive allosteric modulation is a promising area of research to bring about symptomatic as well as potentially disease-modifying treatment.

Another potential specific target for the management of PD is mGluR7s. While mGluR4 is highly expressed in the striatum and the GP, it is little or not expressed in SN where mGluR7 is predominant (Corti et al. 2002). mGluR7s have also high expression in the striatum and very low pallidal expression. In the striatum, mGluR7s are localized presynaptically on the corticostriatal glutamatergic synapses, where they are found to decrease the glutamatergic tone and attenuate transmission in cholinergic interneurons. AMN082 is an mGluR7 selective PAM (Mitsukawa et al. 2005). Intrastriatal injection of this compound reverses haloperidol- and 6-OHDA-induced parkinsonism in rats and haloperidol-induced Parkinsonism in mice. This effect was absent in mGluR7 knockout mice, demonstrating that it is mediated by mGluR7 (Greco et al. 2010). The anti-cataleptic property of the compound was evident only in narrow doses, causing motor impairment at higher doses. AMN082, at high doses, decreases locomotor activity of mGluR7 knockout and wild-type mice, demonstrating that this negative effect is not mGluR7 related (Palucha et al. 2007; Salling et al. 2008).

Apart from the striatum and GP, the SN is another key location for group III mGluRs. Injection of L-SOP or L-AP4 into the SNpr reversed akinesia in the reserpine-treated rat model, while infusion of L-AP4 in the SNpc prevented the akinesia and the nigrostriatal degeneration in the 6-OHDA-lesioned rat. These responses were each inhibited by pretreatment with the group III mGluRs antagonist CPPG, confirming they were mediated by activation of group III mGluRs (Austin et al. 2010). This effect seems to be mediated by presynaptic inhibition of glutamatergic input from the STN. A similar study explored selective targeting of mGluR4, mGluR7, and mGluR8 in the SNpr. Activation of mGluR4 (with PHCCC in combination with a subthreshold concentration of L-AP4) or mGluR7 (with AMN082 also in combination with a subthreshold concentration of L-AP4) reduced the release of the glutamate analogue, [<sup>3</sup>H]-D-aspartate, from SNpr tissue prisms in vitro. In contrast, the mGluR8 selective agonist, (S)-3,4-DCPG, failed to provoke the same inhibition. Provided that glutamatergic afferents from the STN comprise the major excitatory input into the SNpr, mGluR4, and mGluR7,

activation may be effective for restricting glutamate release in this location (Broadstock et al. 2012). Reversal of reserpine-induced akinesia was produced following direct injection of PHCCC or AMN082 into the SNpr. Both of them acted in the absence of a subthreshold dose of L-AP4. The effects of both PHCCC and AMN082 were inhibited by CPPG, the selective group III mGluRs antagonist, confirming that these responses are mediated through group III mGluRs. Contrary to what it should be expected, injection of (S)-3,4-DCPG into the SNpr produced a dose-dependent reversal of akinesia. However, pretreatment with CPPG did not inhibit this effect, suggesting it was driven through other mechanisms independent of group III mGluRs.

Group III mGluRs have also a neuroprotective effect by preventing excitotoxicity induced by iGluRs. In this instance, excess of extracellular glutamate activates presynaptic group III mGluRs, this inhibits glutamate release, and then the activation of postsynaptic NMDARs is diminished (Vera and Tapia 2012).

#### 5 Conclusion

In spite of the innumerable preclinical studies conducted in experimental animal models of PD associated with LIDs, such as rodents and nonhuman primates, some of which generated impressive results, the possibility to consistently attenuate or mitigate the development of LIDs in the clinical setting is still an unmet need. This is mostly due to the incomplete understanding of the neural and molecular substrates of this phenomenon, although numerous studies suggest a decisive role of glutamate receptors on this phenomenon. In fact, non-dopaminergic neurotransmitter receptors expressed by striatal neurons, such as specific subtypes of ionotropic (NMDA) and metabotropic (mGluR5) glutamate receptors, play a crucial role in the modulation of corticostriatal inputs, as well as in the modulation of the striatal output to downstream nuclei of the basal ganglia circuit. As consequence, in addition to the changes directly caused by PD, the involvement of these receptors triggers further modifications in basal ganglia functional organization, which set the ground for the functional alterations that underlie LIDs. Based on these concepts, drugs acting not only directly on glutamate receptors as NMDA and mGluR5 but even indirectly on adenosine A2A receptors have been proposed as antidyskinetic agents, showing auspicious effects. Considering that mGluR5 receptors contribute to these receptor mosaics, it is likely that new anti-LID pharmacological strategies will have to rely on compounds capable of acting, simultaneously, on all the components of these receptor mosaics, thereby combining the effects of adenosine and metabotropic receptor antagonists. However, the ubiquitous location of the different subtypes of glutamatergic receptors leaves open new ways of research in the field of PD.

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# Lead and Excitotoxicity

# Abdur Rahman

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#### Abstract

Lead (Pb<sup>2+</sup>) is a known neurotoxicant, but the mechanism of its neurotoxicity is not clearly understood. Several biochemical alterations have been shown to be caused by Pb<sup>2+</sup> exposure in mammalian brain, but none of these changes alone can explain the mechanism of Pb<sup>2+</sup>-induced impairment of learning and memory. The most mechanistically relevant biochemical abnormalities that are directly involved in learning and memory are the excitotoxic effects caused by modulation of the *N*-methyl-D-aspartate-type glutamate receptors (NMDAR) in glutamatergic synapses. Pb<sup>2+</sup> is known to affect not only the expression of the different subunits of the NMDARs but also the ontogenic developmental switch

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A. Rahman

Department of Food Science and Nutrition, College for Women, Kuwait University, Safat, Kuwait e-mail: abdurrahman.ahmad@ku.edu.kw; abdur\_rahman31@hotmail.com

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of the various NMDAR subunits that is essential for learning and memory. Overactivation of serine/threonine protein phosphatases (PPs) appears to be involved in these synaptic changes. PPs may not only affect the functions of the various subunits of the NMDAR directly by modulating the phosphorylation state of these subunits but may also affect their downstream function by modulation of the phosphorylation state of the downstream effectors like the cyclic AMP response element binding protein (CREB) and other proteins involved in this process. There is a great need to put these isolated pieces of information together and workout the exact pathway(s) that are disturbed by Pb<sup>2+</sup>.

#### Keywords

Excitotoxicity • Hippocampus • Lead • Learning and memory • Neurotoxicity • NMDAR • Protein phosphatases

#### 1 Introduction

Lead ( $Pb^{2+}$ ) is a toxic heavy metal with no known physiological functions in the body. Because of its excessive use in industry, humans have been, and are constantly being, exposed to  $Pb^{2+}$ . Although  $Pb^{2+}$  poisoning in both its acute and chronic forms has gradually declined with a reduction in the use of leaded gasoline over the last few decades and with increasing surveillance of industrial and domestic exposure, it has not disappeared (Dorea 2004; Bellingr and Bellinger 2006).  $Pb^{2+}$  gets into the body through food and water, environmental pollution, agricultural technology, and food processing. Absorption and retention in the body depends on age, chemical environment of the gastrointestinal tract, and nutritional status of the individual. Generally conditions that favor calcium absorption also favor  $Pb^{2+}$  absorption and retention. The total body amount does not affect absorption, as there is no feedback mechanism for its absorption (DeMichele 1984).

Pb<sup>2+</sup> toxicity affects several organ systems including the nervous, hemopoietic, renal, reproductive, endocrine, and skeletal systems (Goyer 1995; Borja-Aburto et al. 1999). The severity of these toxic effects depends on the duration of exposure, dose, and the developmental stage of the subjects. Children are particularly at increased risk of toxicity because of their frequent exposure and increased absorption and retention capacity. The hemopoietic, renal, and reproductive systems are affected at relatively high doses, whereas the central nervous system (CNS) is affected by low doses (Borja-Aburto et al. 1999).

 $Pb^{2+}$  has long been known as a neurotoxic metal but the underlying biochemical mechanism of the neurotoxic effects of  $Pb^{2+}$  is not well understood at present despite the large body of research done on this subject. Three very exhaustive reviews on the health effects of Pb toxicity and in particular on the neurotoxic effects of  $Pb^{2+}$  have been written in the past decade (Toscano and Guilarte 2005; White et al. 2007; Neal and Guilarte 2010). Of the many biochemical changes

studied, the most relevant and best studied area is the excitatory effect of  $Pb^{2+}$  at the glutamatergic synapses. This chapter is aimed to focus on the excitotoxic effects of  $Pb^{2+}$  and to augment these reviews by adding recent findings reported in the literature in the area of  $Pb^{2+}$  neurotoxicity.

#### 2 Neurotoxicity of Lead

Pb<sup>2+</sup> neurotoxicity is known since the late nineteenth century (reviewed by Toscano and Guilarte 2005). While high blood Pb<sup>2+</sup> levels (>70 µg/dL) are known to cause overt encephalopathy (Chisolm 2001), low levels (as low as 7 µg/dL) in children are associated with neurobehavioral and endocrine alterations such as an increase in hyperactivity and distractibility, delayed puberty, and cognitive deficits in the form of IQ changes (Needleman et al. 1979; Bellinger 1995; Nevin 2000; Selevan et al. 2003). Studies have demonstrated a cognitive deficit of 0–5 points on the IQ scale for every 10 µg/dL increase in blood Pb level (Bellinger 1995). It has been estimated by the USEPA that a reduction of a single IQ point results in an added financial burden of US\$ 8,346 per intoxicated child (Reviewed by Toscano and Guilarte 2005).

The current limit of concern for childhood Pb<sup>2+</sup> intoxication set by the Center for Disease Control and Prevention (CDC) is 10 µg/dL. However, recent data suggest that CNS toxicity may be exerted by blood Pb levels of <10 µg/dL (Lanphear et al. 2000; Canfield et al. 2003), as the nervous system is especially sensitive to the effects of Pb. In a prospective study, a lifetime blood Pb<sup>2+</sup> levels <10 µg/dL was shown to be associated with cognitive deficits in children at 3 and 5 years of age (Canfield et al. 2003). In experimental animal models, Pb<sup>2+</sup>-exposure causes deficits in spatial learning and long-term potentiation (LTP) (Altmann et al. 1991; Kuhlmann et al. 1997; Gilbert and Mack 1998; Nihei et al. 2000). It has been suggested that there is a "developmental window" that spans from gestation through lactation, in which exposure to low levels of Pb<sup>2+</sup> is able to cause long-lasting cognitive function deficits (Kuhlmann et al. 1997). This concept of developmental window has been demonstrated by the reports that children that were previously, but not currently, exposed to Pb<sup>2+</sup> exhibited lasting neurobehavioral and cognitive deficits (Needleman et al. 1979).

The hippocampus plays a pivotal role in learning and memory processes, and it has been suggested that this structure is particularly affected by  $Pb^{2+}$  (Bielarczyk et al. 1996; Sharifi et al. 2002). The most troubling aspect of  $Pb^{2+}$  toxicity in children is that neurotoxicity caused by  $Pb^{2+}$  exposure is irreversible. Chelation therapy, which is the primary means of treating children with blood  $Pb^{2+}$  levels of >45 µg/dL, can reduce the body burden of  $Pb^{2+}$  but does not reverse the cognitive or behavioral deficits associated with  $Pb^{2+}$  exposure (US, CDC 2002; Rogan et al. 2001; Dietrich et al. 2004). This highlights the possibility that  $Pb^{2+}$  exposure induces long-lasting (or permanent) changes in the brain during a critical period of development in childhood (Neal and Guilarte 2010).

Many biochemical changes have been reported in the brain that may explain the mechanism of Pb<sup>2+</sup> neurotoxicity. Some of the reported biochemical alterations

caused by Pb<sup>2+</sup> exposure in the brain include: altered adenylyl cyclase activity and reduction in the heme-containing enzymes and lower energy metabolism in the developing brain (Clarkson 1987), interference with cell adhesion molecules (Silbergeld 1992), reduced activity of alkaline phosphatase (Antonio and Lert 2000), decreased expression (Nihei et al. 2001) and activity (Xu et al. 2005) of protein kinase C, decreased production of transthyretin and low availability of thyroid hormone to the developing CNS (Zhang et al. 1996), decreased levels of nitric oxide in the hippocampus (Sun et al. 2005), altered neurotransmitter activity in the hippocampus (Reddy et al. 2007; Wang et al. 2007), and altered protein phosphorylation and impairment of the glutamatergic synapse transmission (Neal and Guilarte 2010). None of these biochemical changes alone explains the learning and memory deficits caused by the low-dose  $Pb^{2+}$  exposure. There is a need to put these pieces together and workout the pathway(s) involved in learning and memory that are affected by  $Pb^{2+}$  exposure. The understanding of such pathway(s) is essential to devising any therapeutic/intervention strategies to combat Pb<sup>2+</sup>-induced neurotoxicity.

#### 3 Hippocampal Plasticity, NMDA Receptor, and Learning

Hippocampus is the main brain region involved in the acquisition and consolidation of higher brain function, particularly spatial learning and memory. The disruption of hippocampal function by a variety of methods produces deficits in such brain functions (Izquierdo 1993; McNamara and Skelton 1993). The major cellular mechanism within the hippocampus believed to be responsible for acquisition of new memories is LTP, which is a long-lasting increase in synaptic efficacy following brief periods of stimulation of specific synapses (Malenka and Nicoll 1999; Hashemzadeh-Gargari and Guilarte 1999; Nihei et al. 2000, 2001; Shimizu et al. 2000).

Some forms of LTP in the hippocampus, specifically those induced in Schaffer collateral-CA1 and perforant path-dentate gyrus synapses, are dependent upon *N*-methyl-D-aspartate-type glutamate receptor (NMDAR) activation (Madison et al. 1991; Malenka and Nicoll 1993, 1999; Teyler and DiScenna 1987; Zalutsky and Nicoll 1990). LTP requires presynaptic glutamate release and subsequent activation of the postsynaptic NMDAR (Collingridge and Bliss 1987; Massicotte and Baudry 1991; McNaughton 1993). Disruption of NMDAR function pharmacologically or by deletion of specific NMDAR subunits using gene knock-out techniques are associated with disruption of hippocampal LTP and learning and memory (Gilbert and Mack 1990; Robinson and Reed 1992; Morris et al. 1986; Neal and Guilarte 2010).

## 4 NMDA Receptors and Their Structure

Glutamate receptors are classified into metabotropic and ionotropic subtypes according to whether they exist as G-protein-coupled receptor or as an ion channel.

Both these receptor types mediate the actions of glutamate. Metabotropic glutamate receptors (mGluRs) have been extensively studied in hippocampal physiology. mGluRs are composed of eight isoforms (mGluR1–8), which are classified into groups I–III. Of these mGluR5, which belongs to group I, is primarily postsynaptic and is coupled preferentially to Gq/11 and its downstream effectors. Recent studies have demonstrated that mGluR5 is involved in learning and memory. mGluR5 has been shown to be critically important for both hippocampal synaptic plasticity and hippocampus-based learning and memory (reviewed by Xu et al. 2009a, b). Inhibition of mGluR5 with specific antagonist MPEF impaired the acquisition and consolidation of hippocampus-dependent memory, whereas its activation by specific agonist showed the opposite results (Gasparini et al. 1999).

Ionotropic receptors are further classified based on their selective agonists as NMDA, AMPA, and Kainate receptors (Hassel and Dingledine 2006). These receptors bind with glutamate with different affinities. Of these, NMDAR are the most tightly regulated and the most extensively studied. Activation of the NMDAR plays a central role in brain development, learning, and memory as well as in neurodegenerative diseases (Collingridge and Lester 1989; Ozawa et al. 1998; Scheetz and Constantine-Paton 1994). These receptors are located primarily in the hippocampus but also in the cerebral cortex (Monaghan et al. 1983; Monyer et al. 1994; Moriyoshi et al. 1991) and play an essential role in hippocampus-mediated learning and memory (Morris et al. 1982, 1986).

NMDAR is a tetrameric complex assembled from the ubiquitous NR1 subunit which is an essential component of all NMDAR complexes along with various combinations of NR2 or NR3 subunits. NR1 subunit is a constitutional component of NMDA receptors and is widely expressed throughout the CNS at all ages, whereas NR2A and NR2B are functional components whose expression varies with the developmental stage of the animal and the different regions of the brain (Xu and Rajanna 2006). A single gene encodes NR1, but at least eight splice variants of NR1 subunits (NR1A to NR1H) have been found (Laurie and Seeburg 1994; Zukin and Bennett 1995). These splice variants of NR1 impart different pharmacological characteristics to the NMDAR (Durand et al. 1992). Alternatively spliced exon 5 at the N terminus (N-cassette) encodes for a 21 amino acid sequence. Splicing of exon 5 results in transcripts designated as lacking (NR1-a) or containing (NR1-b) the N-cassette (Zukin and Bennett 1995). Exons 21 and 22 encode for two C-terminus cassettes, C1 and C2, which code for 37 and 38 amino acid sequences, respectively. The individual splicing of the C1 or C2 cassette results in transcripts designated as NR1-2 and NR1-3. The presence or absence of both C-terminus cassettes results in NR1–1 and NR1–4 variants. Deletion of the C2 cassette alters the reading frame and results in the creation of an additional coding region of 22 amino acids, the C2' cassette (Toscano and Guilarte 2005).

Compared to NR1, NR2 and NR3 are more complex and mostly determine the function of NMDAR channel. NR2 subunits are encoded by four distinct genes (NR2A–NR2D), while NR3 subunits are encoded by two distinct genes (NR3A and NR3B). NR2 subunits play positive roles in gating the NMDAR channel, while NR3 subunits play negative roles by forming an unconventional channel

(Perez-Otano and Ehlers 2004). The molecular biology of these subunits, their deferential developmental and regional expression and their distinct intracellular protein associations and functions are thoroughly reviewed by Neal and Guilarte 2010. Together, the associations between NR1 splice variants with other subunits exhibit an exquisite degree of heterogeneity and specialization and play essential roles in synaptic activity (Neal et al. 2011).

NMDAR subunits play critical roles in hippocampal synaptic plasticity. Blockades of NMDAR containing either NR2A or NR2B subunits lead to a selective defect in either LTP or long-term depression (LTD), respectively (Liu et al. 2004). The differential expression of various NMDR subunits across the developmental time span of peak LTP and hippocampus-mediated learning, together with the impairment of learning due to targeted knockout of NMDAR subunits strongly support the essential role of NMDAR in synaptic plasticity and learning and memory (Neal and Guilarte 2010).

## 5 Activity-Dependent Expression of NMDA Receptor Subunit

The expression of synaptic NMDAR subunit is controlled not only by a genetically programmed development of excitatory circuitry but also by the level of activity present at the synapse. Both in vivo and vitro studies from neuronal cultures have provided evidence for the activity-dependent modifications in NMDAR subunit expression (Toscano and Guilarte 2005). The expression of NR2A subunit, but not NR2B, has been shown to be dependent upon calcium influx mediated by NMDAR and L-type calcium channels since pharmacological blockade of NMDAR and L-type calcium channels decreased NR2A subunit expression but had no effect on NR2B expression. This resulted in the expression of NMDAR complexes with higher proportion of NR2B subunits with the corresponding functional implications (Hoffmann et al. 2000). A reduction in presynaptic exocytosis produced similar results (Lindlbauer et al. 1998). In addition, activity-dependent changes of NMDAR subunits also occur at the postsynaptic density (PSD). Pharmacologically blocking of the sodium channels resulted in decreased PSD levels of NR2A but increased levels of NR1 and NR2B subunits, whereas, activating these channels had the opposite effects (Ehlers 2003).

## 6 Lead and Synaptic Transmission

It has been suggested that a number of neurotoxic effects of Pb<sup>2+</sup> may be due to its interference with neurotransmitter systems, particularly those which involve cellular calcium homeostasis, and calcium-dependent enzymes (Guilarte 1997; Finkelstein et al. 1998; Savolainen et al. 1998a, b; Bressler et al. 1999; Goyer and Clarkson 2001; Nihei and Guilarte 2001; Gilbert and Lasley 2002). Establishment of neuronal circuitries in the developing central nervous system depends on the

pattern of electrical activity going through the synapses. At early stages of brain development, most neurons fire spontaneously, and this spontaneous electrical activity is believed to be required for axonal outgrowth, pruning of synaptic connections, and maturation of neuronal signaling properties (Moody 1998). Therefore, it can be inferred that Pb<sup>2+</sup>-induced impairment of learning and memory in children is a result of altered synaptic activity in the brain, particularly in hippocampus which is involved in cognitive processing (Swanson et al. 1997).

The mechanism by which  $Pb^{2+}$  alters synaptic activity remains unknown. It has been proposed that  $Pb^{2+}$  affect synaptic activity by mimicking the activity of calcium (Ca<sup>2+</sup>). The ability of  $Pb^{2+}$  to substitute for Ca<sup>2+</sup> is one of the primary mechanisms proposed for  $Pb^{2+}$  toxicity in the brain. This Ca<sup>2+</sup>-mimetic ability of  $Pb^{2+}$  has been reported to enhance not only spontaneous neurotransmitter release but also inhibit evoked neurotransmitter release due to impeding Ca<sup>2+</sup> influx through voltage-sensitive Ca<sup>2+</sup> channels (Minnema et al. 1988; Kober and Cooper 1976; Atchison and Narahashi 1984; Braga et al. 1999a; Peng et al. 2002; Xiao et al. 2006). Pb<sup>2+</sup> is also known to bind with intracellular Ca<sup>2+</sup>-binding proteins and may prevent the detection of Ca<sup>2+</sup> signaling essential to neurotransmission (Bouton et al. 2001; Marchetti 2003). Ca<sup>2+</sup>-regulated proteins that are known to be targeted by Pb<sup>2+</sup> include protein kinases, protein phosphatases, voltage-gated Ca<sup>2+</sup> channels (VGCC), and Ca<sup>2+</sup>-binding proteins that regulate mobilization and docking of synaptic vesicles (Atchison 2003; Suszkiw 2004; Braga et al. 1999a, b).

 $Pb^{2+}$  is also known to affect the neurotransmission of glutamate, which is the most abundant excitatory neurotransmitter in the brain. Glutamate and its receptors have an important role in LTP and synaptic plasticity, which are fundamental processes involved in learning and memory (Malenka and Nicoll 1999). Chronic developmental  $Pb^{2+}$  exposure of rats as well as in vitro acute  $Pb^{2+}$  exposure of hippocampal cultures and slices have been reported to decrease the release of glutamate and  $\gamma$ -aminobutyric acid (GABA) (Lasley and Gilbert 2002; Braga et al. 1999a; Xiao et al. 2006).

## 7 Lead and LTP

The effects of  $Pb^{2+}$  on hippocampal LTP has been demonstrated since the early 1990s. Developmental  $Pb^{2+}$  exposure increases the threshold for hippocampal LTP induction, reduces the magnitude of potentiation, and accelerates its decay (Reviewed by Toscano and Guilarte 2005; Altmann et al. 1994; Gilbert and Mack 1998; Gilbert et al. 1999a, b). Animals exposed to  $Pb^{2+}$  from early life showed deficient excitatory postsynaptic potential (EPSP) as well as cellular (population spike) components of this field response, and these effects persisted to adulthood despite termination of  $Pb^{2+}$  exposure at weaning (Gilbert et al. 1996, 1999a). Similar effects of  $Pb^{2+}$  exposure on LTP were also reported in hippocampal slices from exposed animals (Altmann et al. 1993; Sui et al. 2000). These data indicate that  $Pb^{2+}$  exposure reduced LTP magnitude and thus impaired the efficacy of the cellular mechanisms that support learning in the hippocampus (White et al. 2007).

This  $Pb^{2+}$ -induced deficits in hippocampal LTP has been suggested to be associated with a reduction in Ca<sup>2+</sup>-dependent glutamate release (Lasley and Gilbert 1996; Lasley et al. 2001). However, the induction of LTP in mossy fiber-CA3, which is dependent upon an increase in glutamate release, is not affected by  $Pb^{2+}$  exposure (Kawamura et al. 2004), while the induction of LTP at excitatory synapses that are not dependent on increased glutamate release but are NMDAR dependent (i.e., Schaffer collateral-CA1 LTP or perforant path-dentate gyrus LTP) is impaired in the  $Pb^{2+}$ -exposed brain. These results clearly indicate that  $Pb^{2+}$ -induced deficits in LTP are NMDAR-specific (Toscano and Guilarte 2005).

#### 8 Lead, Calcium, and Glutamate Release

 $Pb^{2+}$  mimics  $Ca^{2+}$  and disrupts cellular calcium homeostasis as well as the role of  $Ca^{2+}$  as an important intracellular second messenger (Simons 1993). Voltagesensitive calcium channels (VSCCs), which regulate  $Ca^{2+}$  influx, are essential for many neuronal processes such as neurotransmitter release (Kobayashi and Mori 1998).  $Pb^{2+}$  may affect neuronal functions in two ways; first by inhibiting the entry of  $Ca^{2+}$  into the cell, as  $Pb^{2+}$  is a very potent inhibitor of VSCCs (Audesirk 1993; Busselberg et al. 1993; Evans et al. 1991; Minnema et al. 1988). Second,  $Pb^{2+}$  itself may enter cells through  $Ca^{2+}$  channels and by mimicking  $Ca^{2+}$  may affect its functions (Reviewed by Loikkanen et al. 2003).

A dose-dependent reduction in glutamate release in animals developmentally exposed to  $Pb^{2+}$  has been reported (Gilbert and Lasley 2007). Chronic  $Pb^{2+}$ exposure beginning in utero or in the early postweaning period and continuing throughout life alters presynaptic release of glutamate in the rat hippocampus. However, this Pb<sup>2+</sup>-induced glutamate release appears to be dose-dependent. At lower Pb levels synaptic release of glutamate was diminished and this effect was reversed at higher level of Pb<sup>2+</sup> exposure (Lasley and Gilbert 2002). This biphasic dose response indicates the presence of more than one mechanisms of Pb<sup>2+</sup> action. The diminished glutamate release at lower Pb<sup>2+</sup> levels appears to be due to the blocking of VSCC by Pb<sup>2+</sup>. The increased glutamate release at higher level of Pb<sup>2+</sup> exposure suggests the diminution of the K<sup>+</sup>-stimulated transmitter response (White et al. 2007). Similar findings were reported from acute exposure of cultured hippocampal neurons to Pb<sup>2+</sup> (Braga et al. 1999a, b). Recent studies have revealed that the loss of proteins involved in vesicular release, namely, synaptophysin and synaptobrevin, is also involved in Pb<sup>2+</sup>-induced modulation of transmitter release (Neal et al. 2010).

In addition, nicotinic modulation of synaptic transmission has also been implicated in Pb<sup>2+</sup>-induced transmitter release (Nihei et al. 2000). Hippocampal neurons exposed to various concentrations of Pb<sup>2+</sup> inhibited somatodendritic  $\alpha_4\beta_2$  nAChRs and  $\alpha_7$  nAChRs and caused substantial inhibition of transmitter release. This effect was found to be due the direct interaction of Pb<sup>2+</sup> with these receptors (Ishihara et al. 1995; Mike et al. 2000). It has also been hypothesized that Pb-induced inhibition of nicotinic cholinergic modulation of action potential-dependent transmitter release is mediated by a PKC-dependent mechanism (Braga et al. 2004). It has been suggested that  $Pb^{2+}$  activates PKC which in turn phosphorylates nAChRs, proteins associated with the receptors, and/or proteins linking the receptors to the action potential-dependent transmitter release process. Direct phosphorylation of nAChRs by PKC has been shown to reduce receptor activity in sympathetic neurons (Downing and Role 1987). However, this hypothesis has been disputed by other investigators (Seguela et al. 1993; Moss et al. 1996; Fenster et al. 1999).

#### 9 Lead and NMDA Receptor

Glutamate receptors, particularly, the NMDAR are one of the most important targets of  $Pb^{2+}$  (reviewed by Xu and Rajanna 2006). The observation that  $Pb^{2+}$ exposure during development affects NMDAR-dependent LTP (LTP dependent on increased glutamate release) but not NMDAR-independent LTP in hippocampus provided the first experimental evidence that Pb<sup>2+</sup> targets NMDARs (Gutowski et al. 1998; Kawamura et al. 2004). Since then, several studies demonstrated that  $Pb^{2+}$  is a selective and potent noncompetitive inhibitor of the NMDAR (Toscano and Guilarte 2005; White et al. 2007; Neal and Guilarte 2010; Neal et al. 2011). Pb<sup>2+</sup> in micromolar concentration causes a reversible inhibition of the current activated by glutamate through the NMDAR channel in cultured and acutely dissociated neurons and reduces access to the NMDA receptor channel in brain tissue homogenates (Marchetti 2003; Lasley and Gilbert 1999). Decreased NMDAspecific glutamate receptor binding has also been reported in the brain of Pb<sup>2+</sup>-exposed rats (Rajanna et al. 1997). A possible mechanism for this  $Pb^{2+}$ -induced inhibition of the NMDAR is the binding of  $Pb^{2+}$  at the zinc (Zn<sup>2+</sup>) regulatory (inhibitory) site of the NMDAR in a voltage-independent manner (Lasley and Gilbert 1999).

The effect of Pb<sup>2+</sup> on NMDA-activated current is dependent on developmental stages and NR subunit types (Ishihara et al. 1995; Omelchenko et al. 1997; Ujihara and Albuquerque 1992). In addition to acting as an NMDAR antagonist, Pb<sup>2+</sup> exposure also disrupts normal NMDAR ontogeny. Pb<sup>2+</sup>-induced changes in NMDAR subunits during development may form the basis for Pb<sup>2+</sup> effects on synaptic plasticity and cognitive function (Reviewed by Lau et al. 2002). Developmental Pb<sup>2+</sup> exposure has been shown to cause alterations in NR1 splice variant expression, NR2 subunit ontogeny, and NMDAR-dependent signaling (Neal et al. 2011). Adult rats exposed to  $Pb^{2+}$  during development and postweaning into adolescence suffered marked reductions in gene expression of the NR1 subunit of the NMDAR in the hippocampus (Monyer et al. 1992). Chronic developmental  $Pb^{2+}$  exposure in animals is known to alter expression of NR1 splice variants (Guilarte and McGlothan 2003; Xy et al. 2002; Guilarte et al. 2000). The mechanism of these Pb<sup>2+</sup>-induced deficits in NR1 is proposed to be the altered targeting and cell surface expression of NMDAR subunits to the synapse due to changes in NR1 splice variant expression. The lower levels of NR1 subunit mRNA expressed in the Pb<sup>2+</sup>-exposed hippocampus are principally due to decreased levels of the NR1-4 and NR1-2 splice variants. A unique characteristic of these splice variants is that they lack the C1 cassette and impart the highest cell surface expression, PKC potentiation, and calcium kinetics to NMDAR complexes (Neal et al. 2011).

NR2A and NR2B are abundantly expressed in the hippocampus and may be involved in mediating Pb<sup>2+</sup>-toxicity (Guilarte and McGlothan 2003; Lau et al. 2002; Marchetti 2003; Nihei and Guilarte 1999, 2001; Nihei et al. 2000; Perez-Otano and Ehlers 2004; Toscano et al. 2002; Waters and Machaalani 2004). Chronic  $Pb^{2+}$  exposure alters the composition of the NR2 subunits of the NMDR in the rat brain (Nihei and Guilarte 1999). Specifically, developmental Pb<sup>2+</sup> exposure in animals decreased expression of the NR2A subunit with no change or a small increase in NR2B subunit expression (Nihei et al. 2000; Nihei and Guilarte 1999; Zhang et al. 2002; Guilarte and McGlothan 1998; Toscano et al. 2002; Neal et al. 2011). Chronic Pb<sup>2+</sup> exposure not only reduces NMDAR level but also prevents or delays the developmental switch of NR1/NR2B complex to NR1/ NR2A complex with reduction of activity-dependent synaptic plasticity in the mature brain. During early development, NR2B-containing NMDARs predominate until a developmental switch occurs, resulting in the incorporation of the NR2A subunit (Monyer et al. 1994). It is suggested that this developmental switch from predominately NR2B-containing NMDARs to NR2A-containing NMDARs is delayed or impaired during Pb<sup>2+</sup> exposure (Toscano et al. 2002; Toscano and Guilarte 2005).

These NR2 subunit-specific effects of lead could be explained by the differential sensitivities of different subunits to  $Pb^{2+}$ . Gavazzo et al. (2008) have shown that  $Pb^{2+}$  interacts at the  $Zn^{2+}$  regulatory site of NMDAR complexes containing the NR2A but not the NR2B subunit. The NR2 subunits have different  $Zn^{2+}$  binding sites. NR2A-NMDAR binds  $Zn^{2+}$  at a high affinity (nM affinity), while the NR2B-NMDAR binds  $Zn^{2+}$  with lower affinity ( $\mu$ M range) (Paoletti et al. 2000; Rachline et al. 2005). These findings have been corroborated by the observations that recombinant NR1/NR2A complexes are more sensitive to  $Pb^{2+}$  inhibition than NR1/NR2B complexes (Omelchenko et al. 1996). Since  $Pb^{2+}$  and  $Zn^{2+}$  have similar potencies in inhibiting the NMDAR (Guilarte et al. 1994, 1995), it is likely that changes observed with NR2A-NMDARs but not NR2B-NMDARs may be a result of preferential inhibition of  $Pb^{2+}$  for NR2A-NMDARs (Guilarte et al. 1995; Gavazzo et al. 2008).

Furthermore, different concentrations of  $Pb^{2+}$  have differential influence on NMDAR subunits, and different subunits of NMDAR display different sensitivities to  $Pb^{2+}$ . A dose-dependent reduction by Pb in NMDAR-NR1, NR2A, and NR2B protein levels has been reported (Lau et al. 2002). The rank order of sensitivity of different subunits to Pb inhibition has been reported as follows: NR1A-NR2B > NR1A-NR2A > NR1A-NR2C (Gavazzo et al. 2001) and NR1B-NR2A > NR1B-NR2C > NR1B-NR2D > NR1B-NR2AC (Omelchenko et al. 1997).

Region-specific effects of  $Pb^{2+}$  on the expression of NR1 and NR2B subunits have also been reported. In cortical neurons, expression of NR1 was unchanged but that of NR2B was significantly increased by  $Pb^{2+}$ . In contrast, expression of

both NR1 and NR2B was significantly decreased in hippocampal neurons. Thus, it is likely that the toxic effects of  $Pb^{2+}$  may cause differential damage to different types of memory that are mediated by cortical and hippocampal neurons, respectively (Lau et al. 2002). Furthermore, both regional and developmental differences in the hippocampal neurons have been reported in Pb<sup>2+</sup>exposed rats. For example, the expression of NR1-2a mRNA in Pb<sup>2+</sup>-exposed rats was significantly increased in areas CA1, CA4, and dentate gyrus (DG) at postnatal days (PND) 14-15 but in areas CA4 and DG at PND20-21 (Zhang et al. 2002; Guilarte and McGlothan 1998; Guilarte et al. 2000). On the other hand, Pb<sup>2+</sup>-induced decreased expression of NR2A was observed in areas CA1, CA2, CA3, and DG at PND15 and areas CA1, CA3, and DG at PND20. Similarly, NR3A mRNA levels were also significantly decreased in CA1. CA4, and DG subfields at PND15 and CA1 and DG subfields at PND20 in Pb<sup>2+</sup>-exposed rats (Zhang et al. 2002). These regional and developmental regulations of NMDAR mRNA splicing may lead to abnormality of natural NMDAR stoichiometry, control sensitivities to phosphorylation, and therefore kinetic properties of the NMDA channels and their involvement in LTP and synaptic plasticity (Zhang et al. 2002). Thus, this disruption of the ontogenetically defined pattern of NMDAR subunit expression and NMDAR-mediated calcium signaling in glutamatergic synapses appears to be the mechanism of Pb<sup>2+</sup>-induced neurotoxicity and deficits in learning and memory. These changes are associated with deficits in long-term potentiation in the hippocampus and impairment of spatial learning (Nihei and Guilarte 2001; Toscano and Guilarte 2005; Zhang et al. 2002; Gilbert et al. 1996; Neal and Guilarte 2010).

#### 10 Lead and NMDA Signaling

It has been reported that synaptic NMDARs mediate cyclic AMP response element binding protein (CREB) and extracellular signal-regulated kinases (ERK) activation, synaptic plasticity, and survival pathways. On the other hand, extra-synaptic receptors are associated with a CREB-shutoff pathway, ERK inactivation, and induction of cellular death pathways (Hardingham et al. 2002; Vanhoutte and Bading 2003; Ivanov et al. 2006). Changes in NMDAR subunit composition can result in altered NMDAR-dependent signaling. Many signaling pathways are dependent on NMDAR subunit composition and/or localization. Pb<sup>2+</sup>-induced alterations in NMDAR subunit composition could interfere with the downstream signaling pathways including MAPK signaling (Cordova et al. 2004), calcium/ calmodulin kinase II (CaMKII) activity (Toscano et al. 2005), protein kinase C (PKC) activity (Bressler et al. 1999), and CREB phosphorylation status and binding affinity (Toscano et al. 2002; 2003). CREB is a transcription factor for many NMDAR activity-dependent immediate early genes (IEGs), which play an essential role in memory consolidation (Athos et al. 2002; Bourtchuladze et al. 1994). Thus, CREB plays an important role in signal propagation from synapses to the nucleus by linking NMDAR activation and calcium-dependent signaling to the expression of genes necessary for synaptic plasticity (Bredt et al. 1992). Altered IEG expression in animals exposed to  $Pb^{2+}$  has been observed (Kim et al. 2002), indicating that altered CREB activity due to  $Pb^{2+}$ -mediated disruption of NMDAR signaling may result in impaired learning and memory processes.

CREB phosphorylation is an important element of LTP. Induction of LTP (and learning) increases CREB phosphorylation (Mizuno et al. 2002; Schultz et al. 1999; Viola et al. 2000), whereas pharmacological inhibition of NMDAR-mediated signaling decreases CREB phosphorylation (Athos et al. 2002). CREB phosphorylation at serine-133 facilitates the recruitment of CREB-binding proteins and the assembly of transcriptionally active complex at the start site of CRE containing genes (Chirivia et al. 1993). Pb<sup>2+</sup> exposure decreases CREB phosphorylation without affecting the total CREB levels and alters the ability of CREB family proteins to bind with CRE (Toscano et al. 2003), suggesting that the mechanisms that regulate the phosphorylation of CREB at serine-133 and the binding ability of CREB family proteins may be altered by Pb<sup>2+</sup> exposure. This may affect the transcription of genes associated with learning, memory, and synaptic plasticity.

## 11 Lead and BDNF Signaling

Another potential signaling pathway that may be involved in Pb<sup>2+</sup>-induced neurotoxicity is impaired NMDAR-dependent retrograde signaling of neurotrophic factors, particularly of brain-derived neurotrophic factor (BDNF). Pb<sup>2+</sup>-exposed hippocampal neurons exhibit reduced proBDNF expression and BDNF release. Furthermore, complete recovery of Pb<sup>2+</sup>-induced changes in presynaptic protein levels and vesicular neurotransmitter release has been reported in hippocampal neurons incubated with exogenous BDNF (Neal et al. 2010). Retrograde BDNF signal from the postsynaptic side has been implicated in axon morphology, synaptic connectivity, and synaptic ultrastructure (Neal and Guilarte 2010; Neal et al. 2010). NMDAR activation results in the generation and release of BDNF (Hartmann et al. 2001; Jiang et al. 2005; Walz et al. 2006), which may be essential to the generation or unmasking of presynaptic neurotransmitter release sites (Walz et al. 2006). BDNF signaling is known to modulate the expression of several pre- and postsynaptic proteins (Pozzo-Miller et al. 1999; Tartaglia et al. 2001). Of particular interest is the enhancement of NR2A expression but not the expression of NR2B in hippocampal slices exposed to exogenous BDNF (Small et al. 1998; Caldeira et al. 2007). These results are further supported by the reduced expression of NR2A but not the NR2B subunit and reduced vesicular release in BDNF knockout mice (Margottil and Domenici 2003). These observations suggest that NR2A-NMDARs may be preferentially linked to BDNF signaling. The altered expression of NR2A subunit by both Pb<sup>2+</sup> exposure and impaired BDNF signaling strongly supports the hypothesis that the disruption of NMDAR activity-dependent BDNF signaling in involved in Pb<sup>2+</sup>-induced toxicity in synapses (Neal et al. 2010).

## 12 Lead and Other Brain Cells

Most of the work on Pb<sup>2+</sup>-induced neurotoxicity has focused on neuronal cells. Other support cells, which are severalfold greater in number than neurons and regulate/modulate the function of neurons, may also be affected by Pb<sup>2+</sup> and thus may be involved in Pb-induced neurotoxicity. Of the support cells, astroglia are capable of uptaking 14-fold more  $Pb^{2+}$  than neurons. The uptake of  $Pb^{2+}$  by astroglia is suggested to be through voltage-dependent Ca<sup>2+</sup> channels (Simons and Pocock 1987). This Ca<sup>2+</sup>-dependent uptake of Pb<sup>2+</sup> by astroglia may be induced by their interaction with neurons (Tiffany-Castiglioni 1993; Lindhal et al. 1999) and may involve glutamate-dependent increase in intracellular calcium (Cornell-Bell et al. 1990). The accumulation and storage of  $Pb^{2+}$  in astroglia may be a mechanism of protection for neurons which are more sensitive than astroglia to the toxic effects of  $Pb^{2+}$  (Holtzman et al. 1987: Tiffany-Castiglioni et al. 1986; Tiffany-Castiglioni 1993). On the other hand, such storage of  $Pb^{2+}$  in astroglia may provide a reservoir for its continuous release and thereby may contribute to the toxicity of adjacent neurons or astroglia themselves (Struzynska 2009).

## 13 Lead and Metabotropic Glutamate Receptors

Of the metabotropic glutamate receptor, mGluR5 has also been linked to Pb<sup>2+</sup>-induced deficits in learning and memory. The impact of developmental lead exposure on hippocampal mGluR5 expression and its potential role in Pb<sup>2+</sup>-induced neurotoxicity has been investigated both in vitro and in vivo (Xu et al. 2009a, b). Decreased expression of mGluR5 mRNA and protein by Pb<sup>2+</sup> dose dependently suggests that mGluR5 might be involved in Pb<sup>2+</sup>-induced neurotoxicity. Impairment of mGluR5-dependent long-term depression (LTD) (Huang and Hsu 2006) and decreasing NMDAR-dependent or protein synthesis-dependent long-term potentiation have been hypothesized as potential mechanisms of Pb<sup>2+</sup>-induced neurotoxicity (Toscano and Guilarte 2005; Topolnik et al. 2006; Manahan-Vaughan et al. 2003; Manahan-Vaughan and Braunewell 2005; Naie and Manahan-Vaughan 2004; Neyman and Manahan-Vaughan 2008).

## 14 Protein Phosphatases and Lead-Induced NMDAR-Dependent Neurotoxicity

The major Ser/Thr protein phosphatases (PPs) in mammalian brain are PP1, PP2A, and PP2B (Liu et al. 2005). Overactivation/overexpression of these Ser/Thr PPs has been implicated in impairment of learning and memory. PPs are strong molecular constraints on learning and memory (Lee and Silva 2009). PPs are also involved in cognitive decline in aging (Mansuy and Shenolikar 2006; Knobloch et al. 2007) and in favoring forgetting (Genoux et al. 2002). Overactivation of PP1 is associated

with learning and memory impairment (Genoux et al. 2002, 2011; Koshibu et al. 2009, 2011; Graff et al. 2010; Haege et al. 2010). PP2A and PP2B also adversely affect learning and memory (Bennett et al. 2003; Havekes et al. 2006; Yamashita et al. 2006; Mauna et al. 2010; Oberbeck et al. 2010). Overactivation of PP2A induces LTD in vivo (Thiels et al. 1998), and inhibition of PP2A abrogates LTD induction (Mauna et al. 2010).

Whether the overactivation/overexpression of these PPs is involved in Pb<sup>2+</sup>induced impairment of learning and memory has largely remained uninvestigated until now. This issue was recently addressed by investigating the effect of Pb<sup>2+</sup>exposure on PPs both in cultured human neurons and in rats exposed to Pb<sup>2+</sup> during early development. In the Pb<sup>2+</sup>-exposed cultured human fetal neurons, a significant increase in total PPs and PP2A activities was observed. Pb<sup>2+</sup> exposure significantly increased the expression of PP1 and PP2B but significantly decreased the expression of PP2A and PP5 in cultured human fetal neurons (Rahman et al. 2010). In rats exposed to Pb<sup>2+</sup> during early development, learning, short-term memory (STM), and long-term memory (LTM) were significantly impaired at PND 21, and this impairment of learning and LTM was associated with decreased synaptogenesis and increased expression and activity of PP1 and PP2A in the hippocampus. On the other hand, at PND 30, learning and short-term memory (STM), but not the LTM, was impaired by Pb<sup>2+</sup>. At this developmental stage, expression and activity of hippocampal PP1 was increased but that of PP2A was decreased. These results suggest that increased PP1 activity in hippocampus is involved in the impairment of learning and LTM, whereas, increased PP2A activity is involved in the impairment of STM (Rahman et al. 2012).

LTM involves protein synthesis, growth, and formation of new synapses (Martin et al. 1997; Ma et al. 1999; Toni et al. 1999; Bozdagi et al. 2000; De Roo et al. 2008). By contrast, STM involves covalent modification of proteins in the presynaptic or postsynaptic structures (Kandel 2001; Malinow and Malenka 2002). The major covalent modification underlying these long-lasting changes in synaptic communication is protein phosphorylation, which is regulated by a balance between protein kinases (PKs) and PPs. Increased activity of PKs results in increased protein phosphorylation which has been implicated in LTP. On the other hand, increased activity of PPs results in decreased protein phosphorylation, which has been implicated in LTD (Roberson et al. 1996; Winder and Sweatt 2001; Blitzer et al. 2005). Reversible protein phosphorylation regulates presynaptic and postsynaptic events in excitatory and inhibitory neurons. The major substrates for PPs at these synaptic sites include ligand-gated ion channels and G-protein-coupled receptors, whose functional properties, trafficking, and synaptic organization are controlled by reversible phosphorylation (Swope et al. 1999).

Several mechanisms have been proposed to explain the role of PPs in learning and memory. These include dephosphorylation and the subsequent deacetylation of histone proteins and chromatin remodelling and altered expression of CREB and NF- $\kappa$ B (Kandel 2001; Fischer et al. 2007; Miller et al. 2008; Koshibu et al. 2009, 2011; Oberbeck et al. 2010). Both PP1 and PP2A dephosphorylate CREB (Mauna et al. 2010) and thereby reduce CREB-mediated gene expression (Wadzinski et al. 1993; Alberts et al. 1994; Genoux et al. 2002; Oberbeck et al. 2010). Other mechanisms include dephosphorylation by PP1 and PP2A of NMDA receptor and MAPK (Chan and Sucher 2001; Oberbeck et al. 2010), calcium-/ calmodulin-dependent protein kinase IV (CaMKIV) (Westphal et al. 1998; Anderson et al. 2004), ERK (Davis et al. 2000; Norman et al. 2000; Silverstein et al. 2002; Ho et al. 2007), and the AMPA receptor (Thiels et al. 2002).

Of particular significance in the Pb<sup>2+</sup>-induced excitotoxicity is the dephosphorylation and subsequent inhibition of the NMDARs-associated signaling pathways. Protein phosphorylation has been established as an important mechanism for the regulation of NMDAR function. LTP is accompanied by increased glutamate receptor phosphorylation through various protein kinases and a concomitant decrease in protein phosphatase activity (Bliss and Collingridge 1993; Mulkey et al. 1993; Soderling and Derkach 2000). In contrast, LTD has been shown to be dependent on glutamate receptor dephosphorylation mediated by an increase in the activity of protein phosphatases, possibly PP1 and PP2A (Lee et al. 1998, 2000; Thiels et al. 1998). Thus, coordination of kinase and phosphatase activities is crucial for the modulation of synaptic plasticity. Both kinases (PKA) and phosphatases (PP1 and PP2A) are located in physical proximity to NMDAR (Chan and Sucher 2001). Synaptic NMDA receptors are phosphorylated or dephosphorylated depending on synaptic stimulation. Phosphorylated NMDA receptors have enhanced channel openings and a consequent increase in Ca<sup>2+</sup> influx, and this influx is necessary for inducing long-term neuronal changes (Levitan 1999; Prybylowski and Wenthold 2004; Raymond et al. 1994; Roche et al. 1994). Following excitatory neurotransmission and Ca<sup>2+</sup> influx, NMDA receptors are phosphorylated by PKA then rapidly dephosphorylated by PP1, PP2A, or PP2B. This reversible phosphorylation controls synaptic strength, memory formation, and storage by the induction of LTP or LTD (Blitzer et al. 2005). Several PKs, including CaMKII, PKC, PKA, and ERKs, have been implicated in LTP induction and maintenance, whereas PPs are implicated in the induction of LTD. Previous studies have shown that LTD is dependent on the change of the phosphorylation state of glutamate receptors in general (Lee et al. 1998, 2000) and the activity of PPs, particularly PP1 and PP2A (Mulkey et al. 1993, 1994; Thiels et al. 1998). Stimulation of NMDARs activates PP1, PP2A, and PP2B whose substrate specificity is primarily determined through interaction with regulatory and targeting proteins (Mansuy and Shenolikar 2006).

NMDAR activity is attenuated by PP1, PP2A (Wang et al. 1994), and PP2B (Mulkey et al. 1994; Wang and Kelly 1997). PP1 diminishes NMDA-receptormediated synaptic currents in an activity-dependent manner in the hippocampus (Wang et al. 1994; Westphal et al. 1999). PP1 and PP2A appear to decrease the single channel open time in cultured rat hippocampal neurons and thus reduce NMDAR activity (Wang et al. 1994). PP1 has been reported to have more involvement than PP2A in dephosphorylating the NMDAR.

Both positive and negative effects of PP2B on glutamate receptor functions have been reported. PP2B reduces NMDA-receptor-mediated currents by

dephosphorylating the NR2A subunit of the NMDAR and reduction of the open time of individual channels (Lieberman and Mody 1994) and desensitization of the receptor (Krupp et al. 2002; Tong et al. 1995). On the other hand, PP2B has been reported to potentiate NMDAR-induced metabotropic mGluR5 receptor activity by dephosphorylating PKC-dependent sites on the mGluR5 C-terminus and attenuating mGluR5 desensitization (Alagarsamy et al. 2005). PP2B-mediated dephosphorylation also contributes to the developmental switch of NMDAR subunits and thus modulates NMDA receptor subunit composition. In the developing cerebellum, it counteracts the TrkB- and ERK1/2-dependent upregulation of the NR2C subunit, which then exchanges with NR2B to promote synaptic transmission from mature mossy fibers onto granule cells (Suzuki et al. 2005).

The role of PP2A in NMDAR regulation and functioning has been well-studied and best understood. There are intriguing functional similarities between PP2A and NMDAR. Both contribute to the regulation of neural functions such as synaptic transmission and plasticity. Immunoprecipitation studies indicated that the carboxyl domain of NR3A subunit of NMDAR forms a stable complex with the catalytic subunit of PP2A in the rat brain in vivo (Chan and Sucher 2001). Further studies demonstrated that NR3A constitutively associates with the PP2A holoenzyme but not the core enzyme in rat brain synaptic plasma membranes. A sequence of six amino acids between leucine 958 and histidine 974 of the NR3A is critical for binding to PP2A, as deletion or mutation of this sequence disrupted the NR3A-PP2A interaction (Ma and Sucher 2004). Association of PP2A with NMDARs leads to an increase in the activity of PP2A and the dephosphorylation of Ser 897 of the NMDAR subunit NR1. Stimulation of NMDARs leads to the dissociation of PP2A from the complex and the reduction of PP2A activity (Chan and Sucher 2001).

The NR1 subunit is phosphorylated by both PKA and PKC (Tingley et al. 1997). PKA is associated with NMDAR through the scaffold protein yotiao (Westphal et al. 1999), and this NMDAR-associated PKA has been shown to phosphorylate the Ser 897 of NR1 (Chan and Sucher 2001). PP2A associated with NR3A dephosphorylate this serine residue of NR1. PP1 has also been shown to be associated with NMDAR through the same scaffold protein. These PP2A- (and possibly, PP1-) induced changes in phosphorylation of NR1 decrease NMDA currents through the attenuated opening of the NMDAR-gated channels (Wang et al. 1994). PP2A may also indirectly modulate the phosphorylation of NR1 subunits by modulating the function of PKC. It has been reported that PP2A dephosphorylates and inactivates PKC in vitro, and inhibition of PP2A by okadaic acid prevents dephosphorylation of PKC and enhances the function of PKC (Hansra et al. 1996; Millward et al. 1999; Ricciarelli and Azzi 1998).

In addition to NR1, PP2A is also known to dephosphorylate NR2B subunit. It is known that CaMKII can bind to the C-terminal region of the NR2B subunit of NMDAR and phosphorylate it at serine-1303 (Mayadevi et al. 2002). PP2A may regulate the phosphorylation of this site of NR2B subunits by modulating the function of CaMKII. CaMKII autophosphorylated at Thr286 is active, whereas the activity is decreased by dephosphorylation of this site (Colbran 2004; Otmakhov et al. 2004). CaMKII autophosphorylated at Thr286 was first reported to be exclusively dephosphorylated by PP1 in isolated synaptosomes (Shields et al. 1985).

From the literature reviewed above, it is evident that our knowledge about the mechanism of  $Pb^{2+}$ -induced neurotoxicity is patchy, and there is a need to put these isolated pieces of information together to understand how the pathway(s) involved in learning and memory are affected by  $Pb^{2+}$ . Only when we understand the problem, we will be able to devise an effective intervention.

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# **MS and Excitotoxicity**

# Chai K. Lim

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#### Abstract

In this section, it will be studied how excitotoxicity contributes to neurodegeneration in the context of multiple sclerosis (MS). MS has long been regarded as an autoimmune inflammatory disease targeting oligodendrocytes of the central nervous system (CNS). However, recent advancement in research questioned whether MS is merely an inflammatory disease despite pathological evidences of degeneration are clearly present. More recent findings further challenge the research focus of MS which previously thought oligodendrocytes as the main target, whereas current debate points towards pathology of MS being primarily dysfunctional glial interaction, i.e., astrocytes and microglia. Herein, a brief overview will be provided on how various aspects of excitotoxicity are evident in MS pathology and its possible roles between interactions of cells in the CNS that ultimately lead to neurodegeneration in inflammatory condition of MS.

C.K. Lim

Department of Pharmacology, School of Medical Sciences, University of New South Wales, Sydney, Australia

MND and Neurodegenerative Diseases Research Group, Macquarie University, Sydney, Australia e-mail: Chai.Lim@unsw.edu.au

### Keywords

Excitotoxicity • Kynurenine pathway • Multiple sclerosis • Neurodegeneration • Quinolinic acid

# 1 Introduction

Multiple sclerosis (MS) is a chronic degenerative disease of the central nervous system (CNS) characterized by the presence of multifocal lesions with immune cells infiltration, inflammation, demyelination, and neural damages. Although the cause of MS is unknown, very often, MS begins with an autoimmune response to myelin components and progressively leads to oligodendrocytes and axonal degeneration in the chronic phase. The early stages of MS are often resembled by the cycle of inflammation described as relapsing-remitting (RR) phase where the recovery (remitting) period can generally occur from several months to years. About half of RRMS patients will experience exacerbation of the disease progression, of which the remitting period progressive (SP) stage. It remains unclear when and how this switch occurs, but the current hypothesis is that the early stage of MS is driven by the adaptive immunity through inflammation, whereas in the later stage of MS, the innate immunity mediating through neurodegeneration predominates.

# 2 MS, an Autoimmune Disease or Neurodegenerative Disease?

MS has long been considered an autoimmune demyelinating disease of the CNS. The concept of immune-mediated response in the pathogenesis of MS is indisputable. Presence of autoreactive antibodies and T-cell-mediated destruction on CNS myelin had been the core of this concept. Studies using immunized animals with myelin-related peptides known as experimental autoimmune encephalomyelitis (EAE) exhibiting similar phenotype to MS further highlight the importance of the immune regulation in MS progression. As a result, large number of transacted axons arises from inflammatory insult, but most of these demyelinated axons initially survive and subsequently the damages are reversed. However, in late stages of MS, i.e., SPMS, the brain undergoes continuous atrophy where new inflammatory demyelinating lesions are rare. Progressive neuronal and axonal loss provided a logical explanation for the transition from RRMS to SPMS and for continuous and irreversible neurological decline in SPMS. However, these observations present a more typical hallmark of neurodegeneration. More importantly, neuronal and axonal degeneration must continue in environments other than the inflammatory demyelinating lesion. Furthermore, the inflammatory microenvironment contains a variety of substances that could injure axons, such as proteolytic enzymes, cytokines, oxidative products, and free radicals produced by activated immune and glial cells. Hence, mechanisms other than inflammatory demyelination of white matter, therefore, must contribute to axonal and neuronal degeneration. It has been well established that excitotoxicity has played a significant role in neurodegeneration, and the next few sections will look at past and recent evidences of excitotoxicity participating in MS pathogenesis.

# 3 Clinical Evidences of Glutamate-Induced Excitotoxicity in MS

It was notable that the concentration in one of the main excitatory amino acids, glutamate, was significantly elevated in the CNS of MS implicated from several clinical studies. Stover et al. showed that the glutamate level in the CSF of active relapsing phase of MS patients was significantly higher than those who are in the silent (remitting) phase of MS (Stover et al. 1997; Sarchielli et al. 2003). This increase was also associated with the severity and progression of the disease (Sarchielli et al. 2003; Barkhatova et al. 1998). More importantly, Srinivasan et al. demonstrated that the increase of glutamate levels was present in acute lesions and normal-appearing white matter but not in chronic lesions of MS patients using in vivo magnetic resonance spectroscopy imaging technique (Srinivasan et al. 2005). In consistent with the clinical data, MS animal model, EAE lesions are also primarily localized in rich glutamatergic area such as cerebellum and the subpial region of the spinal cord (Brown et al. 1982; Tran et al. 1997).

## 4 Focus on Oligodendrocytes

The potential role of glutamate-induced excitotoxicity in MS was observed in experimental data demonstrating the beneficial effect of NMDA and AMPA/ kainate antagonist, memantine in rat EAE showing reduction of axonal and oligodendrocyte damage but do not dampen CNS inflammation (Wallstrom et al. 1996). This indicates that the mechanism of axonal damage and reversible neurological deficit is likely to act through NMDA receptor and independent of autoimmune neuroinflammation. In addition, two independent studies using NBQX, a selective inhibitor of AMPA/kainate, led to significant reduction in clinical impairment with increased survival of oligodendrocytes in EAE (Pitt et al. 2000; Smith et al. 2000). These data further indicate that AMPA/kainate receptor-mediated glutamate excitotoxicity is a chief factor in oligodendrocyte death in EAE. Indeed, oligodendrocytes express functional AMPA/kainate-type glutamate receptor (Holtzclaw et al. 1995; Steinhauser and Gallo 1996), and exposure to low concentration of AMPA, kainate, or glutamate results in death of oligodendrocytes which can be reversed by addition of NBQX (McDonald et al. 1998). Furthermore, the degree of vulnerability in oligodendrocytes to glutamate excitotoxicity is similar to cortical neuron cultures (McDonald et al. 1998). In addition, oligodendrocytes, like astrocytes, also express functional glutamate transporters for glutamate uptake (Domercq et al. 1999) and glutamine synthetase (D'Amelio et al. 1990; Cammer 1990). This further subjects oligodendrocytes to glutamate excitotoxicity should their ability to regulate glutamate be compromised (Domercq et al. 2005).

It was found that GluR1, a calcium-permeable AMPA subunit, was elevated on oligodendrocytes in the borders and within active MS plaques in comparison with white matter reflected by positive GluR1-immuneopositive oligodendrocytes (Newcombe et al. 2008). This upregulation of GluR1 on oligodendrocytes around and within plaques may render these cells vulnerable to glutamate excitotoxic damage and death mediated by intracellular calcium ions. One possible mechanism for the increase in GluR1 may be due to presence of growth factor that is facilitated by recruitment of progenitor neural cells at the lesion sites. Growth factors can induce robust and selective enhancement of GluR1 expression in functional channel of oligodendrocytes (Chew et al. 1997). Hence, growth factor-induced upregulation of GluR1 may increase cell vulnerability to glutamate and indirectly limit remyelination at lesion sites.

## 5 Focus on Microglia and Astrocytes

During inflammation, glutamate is produced and released into the extracellular space by activated leukocytes and microglia in the CNS (Piani et al. 1991; Werner et al. 2001). Astrocytes, on the other hand, are the main glial cells capable of catabolizing glutamate within the CNS (Schousboe et al. 1993) and are thus strategically situated in the vicinity of neurons and oligodendrocytes to prevent damage caused by variation in glutamate extracellular level. Astrocytes express two critical glutamate transporters, GLAST and GLT-1, to maintain glutamate homeostasis in the CNS (Rothstein et al. 1996). In the course of EAE, although high astrocytic reaction is observed reflected by increased GFAP (marker for astrocytes), there is a decrease in expression of astrocytic glutamate catabolic enzymes, glutamine synthetase, and glutamate dehydrogenase, suggesting reduction in the capacity of astrocytes to break down glutamate (Hardin-Pouzet et al. 1997). Further, Ohgoh et al. also found significant downregulation of expression in the protein level of the glial glutamate transporter GLAST (EAAT1) and GLT-1 (EAAT2) in EAE, possibly contributing to increase extracellular glutamate (Ohgoh et al. 2002).

It is inevitable not to mention inflammatory cytokines in regard to microglia as these two entities are closely linked. Inflammatory cytokines can also fuel excitotoxicity in MS. Two proinflammatory cytokines of MS, interleukin (IL)-1 $\beta$ and tumor necrosis factor (TNF)- $\alpha$ , dose-dependently inhibited glutamate uptake in astrocytes, whereas treatment with interferon (IFN)- $\gamma$  alone promotes glutamate uptake. Hu et al. showed that this paradigm can be antagonized by nitric oxide synthase (NOS) blocker, N<sup>G</sup>-monomethyl-L-arginine, suggesting a mechanistic involvement of NO in IL-1 $\beta$ -mediated inhibition of glutamate uptake (Hu et al. 2000). In addition, anti-inflammatory cytokine IFN- $\beta$ , a common treatment therapy used in RRMS, was also found to mimic NOS inhibitor by limiting NO production while promoting glutamate uptake in astrocytes (Hu et al. 2000). This phenomenon may partly explain the beneficial effects of the current IFN- $\beta$  therapy in MS. TNF- $\alpha$ had also shown to disrupt the glutamate homeostasis by promoting glutamate exocytosis in CXCR4-activated astrocytes (Bezzi et al. 2001). Bezzi et al. also showed that glutamate release is further enhanced by presence of activated microglia (Bezzi et al. 2001). TNF- $\alpha$  was also found to inhibit the uptake of glutamate in a dose-dependent fashion by human primary astrocytes; however, the exact mechanism remains to be resolved (Fine et al. 1996). Hence, it is likely that astrocytes-microglia interaction during MS inflammation, under the influences of proinflammatory cytokines, forms an amplification cascade leading to abnormal glutamate homeostasis. Indeed, Takahashi et al. showed that IL-1 $\beta$  and TNF- $\alpha$  are not detrimental in pure oligodendrocyte culture; however, in co-culture with astrocytes and microglia, they evoke apoptosis in oligodendrocyte mediating through glutamate excitotoxicity (Takahashi et al. 2003). Collectively, all the above evidences suggest that loss of ability to modify the local glutamate levels between glial-glial interactions is likely to contribute to neurodegeneration in MS.

## 6 Quinolinic Acid, Another Potential Endogenous Culprit in Excitotoxicity of MS

One of the NMDA receptor agonists that is worth mentioning and progressively received significant attention in the field of excitotoxicity is quinolinic acid (QUIN). This glutamatergic agonist acts preferentially on distinct population of NMDA receptors containing the NR2A and NR2B subunits. QUIN has had described to be the most potent agonist for NMDA receptors among the others such as NMDA, ibotenate, and glutamate (Monaghan and Beaton 1991). QUIN is one of the downstream metabolites of the kynurenine pathway (KP) which is a major catabolic route of the tryptophan metabolism. It is one of the several metabolites in the KP that are both neuroactive and neurotoxic. Under physiological condition, QUIN is present at nanomolar concentrations in human and rodent brain (Chen and Guillemin 2009). However, during inflammatory state, activation of the KP under the influence of proinflammatory cytokines in the presence of macrophages and microglia can lead to production of up to submicromolar concentration of QUIN (Guillemin et al. 2003, 2005). Hence, its endogenous level in the CNS may be of concern for excitotoxicity given its potency over any other agonists. Indeed, Schwarcz et al. showed that nanomolar concentrations of OUIN injected into rat striatum and hippocampus region of the brain resulted in axon-sparing lesions (Schwarcz et al. 1983). Several in vitro studies also demonstrated that neuronal cells are vulnerable to QUIN excitotoxicity similar to glutamate at pathophysiological concentration including neurons (Kerr et al. 1998), astrocytes (Ting et al. 2009), and oligodendrocytes (Cammer 2001). QUIN can further enhance glutamate excitotoxicity by increasing the release of glutamate by neurons, decrease uptake by astrocytes (Tavares et al. 2002), and inhibit astroglial glutamine synthetase (Ting et al. 2009) consequently disrupting glutamate-glutamine cycle and ultimately leading to excitotoxicity. Detrimental effects of QUIN can be reversed either through increased endogenous of kynurenic acid (KYNA) (Miranda et al. 1997; Harris et al. 1998), an antagonist of the NMDA receptor, or using potent selective NMDA blocker, MK801 and memantine (Braidy et al. 2009). KYNA is also a KP metabolite and its endogenous level in the mammalian system plays a pivotal role in regulating QUIN excitotoxicity. KYNA was found to be present at subnormal concentration in several neurodegenerative diseases (Chen and Guillemin 2009) including MS (Rejdak et al. 2002). It was suggested that it requires twice as much KYNA to antagonist the same concentration of QUIN (Lekieffre et al. 1990). This further suggests that QUIN is a potent NMDA agonist mediating excitotoxicity at pathophysiological conditions such as inflammation. In the context of MS. OUIN had been implicated to be involved in the pathogenesis of MS associated with neurodegeneration (reviewed (Lim et al. 2010)). Flanagan et al. was the first to demonstrate that QUIN is elevated in EAE and is selective to CNS (Flanagan et al. 1995). They further showed that the time course of OUIN production was similar to the neurological signs of the disorder apart from the initial increased in OUIN before onset of any prominent abnormal behavioral sign of EAE. This suggests that QUIN may be involved in the pathogenesis of EAE. Another study reporting similar results further identified the source of OUIN to be coming from infiltrating macrophages resulting in an accumulation of QUIN at a level considered to be toxic to both neurons and glial cells (Chiarugi et al. 2001). In addition, treatment of EAE mice with a KP modulator (Ro61-8048) that selectively inhibits kynurenine monooxygenase (enzyme that breaks down upstream KP metabolite) results in decreased downstream production of QUIN, while upstream metabolite, KYNA, was significantly increased (Chiarugi et al. 2001). Interestingly, this paradigm was associated with reverse paralysis of the EAE which further suggests a detrimental role of QUIN in progression of EAE. These data implied that altered QUIN homeostasis is likely to contribute to MS pathogenesis. However, the level of QUIN in MS patients remains to be elucidated. Clinical data of QUIN involvement in MS may have pharmacological implication for late stages of MS (i.e., PMS) associated with neurodegeneration since there is currently no effective treatment.

## 7 Conclusion

This chapter presented various evidences on the roles of excitotoxicity in a complex multifactorial disease, MS. We have highlighted the various components involved in glutamate excitotoxicity and their interactions in cells and animal model of MS. We also looked at how excitotoxicity may interact with some of the immune components of MS. It is indisputable that all these evidences point towards the pivotal roles of excitotoxicity participating in MS pathogenesis. However, the presence of demyelination, the hallmark of MS, whether it is a cause or effect due to autoimmune response or neurodegeneration remains to be elucidated.

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# Neuroprotectants Targeting NMDA Receptor Signaling

# Sandra M. Vetiska and Michael Tymianski

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### Abstract

*N*-Methyl-D-aspartate (NMDA) receptors are key mediators of fast excitatory synaptic transmission within the mammalian central nervous system (CNS) and play vital roles in learning, memory, and synaptic development. Overactivation of NMDA receptors (NMDARs) at glutamatergic synapses often results in excitotoxicity, which can cause neuronal injury and death, and is associated with many neurological disorders such as ischemic stroke, epilepsy, neuropathic pain, traumatic brain and spinal injuries; ocular disorders such as glaucoma; as well as neurodegenerative diseases, including amyotrophic lateral sclerosis, Parkinson's, Huntington's, and Alzheimer's

e-mail: svetiska@uhnres.utoronto.ca; miketym2@rogers.com

S.M. Vetiska (🖂) • M. Tymianski

Neuroprotection Laboratory, Toronto Western Research Institute, University Health Network, Toronto, ON, Canada

diseases. Numerous NMDAR antagonists have been developed that have been effective in reducing cell damage and death in both in vitro and in vivo experimental models of neurological disease, such as ischemia and traumatic brain injury. Unfortunately, clinical use of these NMDAR antagonists has been limited by intolerable side effects, likely due to the necessity of NMDARs in normal brain function. Intracellular signaling pathways that couple to NMDAR activation can promote either cell survival or cell death, depending on NMDAR subunit composition and/or subcellular localization. Ideal therapeutic strategies targeting NMDAR-mediated neurotoxicity should selectively block pro-death signaling, while sparing pro-survival signals. Neuroprotectants targeting downstream effectors of NMDAR-mediated cell death, rather than NMDARs directly, would maintain normal NMDAR function and minimize adverse effects commonly associated with traditional NMDAR antagonists in the treatment of neurological disorders.

## 1 Introduction

In the mammalian central nervous system (CNS), the principal excitatory neurotransmitter is glutamate. This neurotransmitter acts on two classes of receptors: ionotropic glutamate receptors that form a central ion channel pore and mediate fast excitatory synaptic transmission and metabotropic glutamate receptors, which are G protein-coupled receptors that play neuromodulatory roles within the CNS (Traynelis et al. 2010). Three types of ionotropic glutamate receptors have been identified and classified based on electrophysiological and pharmaco-*N*-Methyl-D-aspartate (NMDA),  $\alpha$ -amino-3-hydroxy-5properties: logical methylisoxazole-4-propionic acid (AMPA), and kainate, each being named for their preferred pharmacologic agonist (Nakanishi 1992; Hollmann and Heinemann 1994). Activation of ionotropic glutamate receptors leads to the opening of their associated ion channel, which, depending on receptor type, is permeable to sodium  $(Na^+)$ , potassium  $(K^+)$ , and calcium  $(Ca^{2+})$  ions. While AMPA and kainate receptors have modest Ca<sup>2+</sup> permeability, NMDA receptors (NMDARs) are highly Ca<sup>2+</sup>-permeable (MacDermott et al. 1986; McBain and Mayer 1994; Dingledine et al. 1999), and it is this  $Ca^{2+}$  influx that is thought to play a major role in both the physiological as well as pathophysiological functions of the NMDAR (Choi 1988a, b; Tymianski et al. 1993; Ghosh and Greenberg 1995).

The term "excitotoxicity," first coined by Olney (1969), refers to the overactivation of glutamate receptors due to excessive synaptic release of glutamate. Initial studies on glutamate toxicity were done by Lucas and Newhouse (1957), who showed that systemic injections of L-glutamate into mice destroy the inner neural layers of the retina. These observations of neuronal death following excessive glutamate administration were later extended to the brain (Olney 1969; Olney and de Gubareff 1978) and spinal cord (Regan and Choi 1991). Excitotoxicity is now considered an important mechanism involved in many acute CNS insults, such as ischemia and traumatic brain injury (TBI) as

well as in chronic neurodegenerative diseases, such as Parkinson's (PD), Huntington's (HD), and Alzheimer's (AD) diseases (Lipton and Rosenberg 1994; Arundine and Tymianski 2004; Lau and Tymianski 2010).

Glutamate-mediated neurotoxicity arises from excessive  $Ca^{2+}$  influx through NMDARs (Berdichevsky et al. 1983; Choi 1987, 1988a, b; Sattler and Tymianski 2000).  $Ca^{2+}$  overload can trigger many detrimental cellular processes, including activation of calcium-dependent proteases, mitochondrial dysfunction, and death signals (Pivovarova and Andrews 2010; Szydlowska and Tymianski 2010). Additionally, excessive cellular  $Ca^{2+}$  can induce the production of harmful reactive oxygen species. For example, nitric oxide (NO), synthesized by the  $Ca^{2+}$ -dependent enzyme neuronal nitric oxide synthase (nNOS), can react with the superoxide anion  $(O_2^-)$  to form the highly reactive species peroxynitrite, which can cause damage to proteins, lipids, and DNA (Radi et al. 1991a, b; Aarts et al. 2003b). NMDARs are key mediators of excitotoxicity in the CNS and have thus been the targets of many neuroprotective strategies.

## 2 NMDA Receptors and Their Antagonists

Structurally, NMDARs exist as heterotetramers, typically consisting of two obligatory NR1 subunits and two regulatory NR2 subunits and less commonly including NR3 subunits (Das et al. 1998; Chatterton et al. 2002; Paoletti and Neyton 2007; Cavara and Hollmann 2008; Pachernegg et al. 2012). Four subtypes of NR2 (NR2A-D) can assemble with the NR1 subunit, most commonly NR2A or NR2B in the forebrain. These receptors are typically diheteromeric in nature; that is, they are comprised of either NR1/NR2A or NR1/NR2B subunits. Some NMDARs can also incorporate two different NR2 subunits (Chazot et al. 1994; Chazot and Stephenson 1997); however, few studies have addressed the functional implications of these receptors.

The type of NR2 subunit determines the biophysical and pharmacological properties of NMDARs; furthermore, these subunits govern distinct protein-protein interactions and downstream signaling pathways, by way of the structural diversity of their carboxyl (C)-terminus (Hardingham and Bading 2003; Cull-Candy and Leszkiewicz 2004; Prybylowski and Wenthold 2004; Kohr 2006; Ryan et al. 2008). Alternatively spliced isoforms also exist for some subunits, such as NR1, creating additional functional diversity (Sugihara et al. 1992; Zukin and Bennett 1995). At resting membrane potential, the channel pore of the NMDAR is blocked by magnesium  $(Mg^{2+})$  which prevents ion flow through the channel. Upon membrane depolarization, Mg<sup>2+</sup> is expelled, allowing for the passage of ions through the channel. Thus, NMDAR activation requires not only the presynaptic release of glutamate but also the depolarization of the postsynaptic membrane. NMDARs also require simultaneous binding of glutamate and glycine (or D-serine) for activation (Johnson and Ascher 1987; Kleckner and Dingledine 1988; Lerma et al. 1990). The glutamatebinding site is located on the NR2 subunits, while the NR1 subunits provide the glycine-binding site (Furukawa and Gouaux 2003; Furukawa et al. 2005).

Evidence from preclinical research has demonstrated a critical role for NMDAR-mediated neurotoxicity in cellular and animal models of neurological disorders, such as ischemic stroke and TBI (Choi 1988a, b; Tymianski et al. 1993; Arundine et al. 2003). NMDA receptor antagonists targeting the glutamatebinding site, the glycine-binding site, or the channel pore, as well as NR2B subunit-selective antagonists, all showed neuroprotection in preclinical studies of excitotoxicity-related CNS diseases (Danysz and Parsons 2002; Wang and Shuaib 2005). Unfortunately, all clinical trials for neuroprotection in acute disorders including stroke and TBI using NMDAR antagonists, such as Selfotel, Aptiganel, and Gavestinel, have failed. This was largely due either to adverse effects or to limited clinical efficacy (Wood and Hawkinson 1997; Morris et al. 1999: Gladstone et al. 2002: Ikonomidou and Turski 2002: Hovte et al. 2004: Muir 2006; Kalia et al. 2008; Lau and Tymianski 2010). In addition to undesirable side effects, a short therapeutic window may be a limiting factor with the use of NMDAR antagonists as neuroprotectants in stroke and TBI (Ikonomidou and Turski 2002; Roesler et al. 2003).

In the case of more chronic disorders, an anti-excitotoxic approach using NMDAR antagonists may have a role. For example, memantine is the only NMDAR antagonist currently approved for the treatment of moderate to severe Alzheimer's disease (AD) in most of Europe, the USA, and Canada, either as a monotherapy or in combination with cholinesterase inhibitors (Thomas and Grossberg 2009). Memantine is a moderate affinity, uncompetitive NMDAR antagonist with strong voltage-dependency and fast kinetics, which allow rapid binding to and quick dissociation from NMDARs. These properties allow memantine to be well tolerated and have few side effects, unlike other NMDAR channel blockers (Danysz and Parsons 2012); however, there is no clear benefit to date for the use of memantine in milder stages of AD and only a small beneficial effect on cognition in vascular dementia (Parsons et al. 1999; McShane et al. 2006; Rammes et al. 2008; Herrmann et al. 2011). Memantine has been studied in several other neurological diseases, both at the preclinical and clinical levels (Stieg et al. 1999; Rao et al. 2001; Volbracht et al. 2006; Rojas et al. 2008; Thomas and Grossberg 2009; Milnerwood et al. 2010), and may have clinical benefit in some disorders, including PD (Aarsland et al. 2009), HD (Beister et al. 2004; Ondo et al. 2007) and HIVassociated dementia (Schifitto et al. 2007; Zhao et al. 2010). Further work is required to establish the full potential of memantine in these diseases.

## 3 Dual Roles of NMDARs in Health and Disease: Implications for Neuroprotectants

### 3.1 Importance of NMDAR Subcellular Localization

The critical roles that NMDARs play in normal brain function may help explain why clinical treatment of stroke and brain trauma with NMDAR antagonists has failed. An ideal therapeutic approach to antagonize NMDAR overactivity in neurological disorders would include blocking downstream excitotoxic signaling without directly blocking NMDARs or inhibiting normal excitatory neurotransmission. Recently, dual roles for the NMDAR in both neuronal death and survival have been increasingly appreciated. These depend, at least in part, on the subcellular NMDAR pool that is activated (Hardingham and Bading 2010). At the neuronal plasma membrane, NMDARs are localized to both the synapse and to extrasynaptic regions, and these two receptor populations may have different roles in physiological and pathophysiological processes. Although NMDARs at both locations can mediate excitotoxicity (Sattler et al. 2000; Liu et al. 2007), studies in cultured cortical and hippocampal neurons showed that stimulation of synaptic NMDARs promotes cell survival pathways, whereas extrasynaptic NMDAR activation is more strongly associated with neuronal death (Vanhoutte and Bading 2003; Soriano and Hardingham 2007; Leveille et al. 2008; Hardingham and Bading 2010). This dual role of NMDARs in cell survival and death stems from distinct intracellular signaling pathways that are coupled to synaptic and extrasynaptic NMDARs. Synaptically localized NMDARs promote cell survival by suppressing apoptotic gene expression (Leveille et al. 2010) and activating pro-survival factors such as cyclic AMP response element-binding protein (CREB), a transcription factor that induces expression of genes such as brain-derived neurotrophic factor (BDNF) that is important for survival (Hardingham et al. 2002). In contrast, extrasynaptic NMDARs can mediate cell death by attenuating the CREB pathway, blocking BDNF expression, and promoting mitochondrial dysfunction and cell death (Hardingham et al. 2002; Gouix et al. 2009). Recent work has demonstrated that, similar to cortical neurons, striatal medium-sized spiny neurons also show increased and decreased CREB signaling after stimulation of synaptic and extrasynaptic NMDARs, respectively (Kaufman et al. 2012).

Alterations in synaptic and extrasynaptic NMDAR localization, leading to an imbalance between pro-survival and proapoptotic signaling, have been associated with several neurological disorders (Gladding and Raymond 2011; Sanz-Clemente et al. 2012). For example, enhanced activity of extrasynaptically localized NMDARs was observed in cerebral ischemia (Tu et al. 2010) and Huntington's disease (HD; Okamoto et al. 2009; Milnerwood et al. 2010). In the striatum of transgenic HD mice, increased NR2B-containing extrasynaptic NMDAR expression, increased current, and associated reductions in nuclear CREB activation were found. Furthermore, this reduction in CREB activity, along with associated motor learning deficits, was reversed by treatment of HD mice with memantine, which was shown to preferentially block extrasynaptic NMDARs (Leveille et al. 2008; Okamoto et al. 2009; Xia et al. 2010). The selective block of extrasynaptic NMDARs by memantine may contribute to its clinical tolerability; that is, memantine may spare physiological synaptic transmission needed for normal brain function but selectively antagonize extrasynaptic NMDARs that are excessively activated under pathological conditions.

## 3.2 Importance of NMDAR NR2 Subunit Subtype

Seemingly contrary to the idea that neuronal survival and death pathways are mediated by synaptic and extrasynaptic NMDARs, respectively, the wellcharacterized synapse-specific protein PSD95 (postsynaptic density-95) is required for NMDAR-mediated excitotoxic neuronal death (Sattler et al. 1999; Aarts et al. 2002; Zhou et al. 2010). This apparent paradox may be explained by the temporally and spatially regulated expression of different NMDAR NR2 subunits (Travnelis et al. 2010). In adult forebrain neurons, NR2A-containing NMDARs are preferentially localized to synapses, while NR2B-containing NMDARs are primarily expressed at extrasynaptic sites (Tovar and Westbrook 1999: Liu et al. 2007: Groc et al. 2009). In addition, NR2A-containing NMDARs are thought to mediate cell survival signals, whereas NR2B-containing NMDARs are associated with cell death pathways, in both in vitro and in vivo models of stroke and TBI (DeRidder et al. 2006; Zhou and Baudry 2006; Liu et al. 2007; Chen et al. 2008; Terasaki et al. 2010). However, this distribution profile of NMDAR NR2 subunits is not absolute. NR2A-containing receptors can be found at extrasynaptic sites; likewise, NR2B-containing NMDARs are expressed in the postsynaptic membrane, where they associate with other postsynaptic density proteins, such as PSD95 (Groc et al. 2006; Thomas et al. 2006; Harris and Pettit 2007). Therefore, neuronal death pathways may be activated by either synaptic or extrasynaptic NR2B-containing NMDARs, while activation of either synaptic or extrasynaptic NR2A-containing NMDARs promotes neuronal survival (Liu et al. 2007).

NMDARs are localized in the cell membrane, including that of synapses, by an array of scaffolding proteins. Among these is the abundant PSD95 protein, discussed in greater detail in sections below. In synapses, PSD95 binds the C-termini of NMDAR NR2 subunits and links NMDARs to other proteins found in the multiprotein complex (MPC) with which NMDARs associate. In doing so, PSD95 links NMDARs to signaling proteins within the MPC that mediate excitotoxic neuronal death (Sattler et al. 1999; Aarts et al. 2002; Lai and Wang 2010; Zhou et al. 2010). The linkage between NMDARs and downstream toxic signaling pathways may be strongest with NR2B-containing NMDARs. Lending support to this idea is a recent study (Martel et al. 2012) that used chimeric constructs of NMDAR NR2A and NR2B subunits with reciprocal exchanges of their C-terminal domains to demonstrate that the NR2B C-terminal domain is important for NMDAR-mediated toxicity, regardless of receptor location. As discussed above in the case of PSD95, the precise mechanisms by which NMDAR subunit composition determines functional outcome of the receptor may lie in the specific intracellular signaling proteins that couple to NMDARs, most often via their C-terminal tails. Ideally, a therapeutic strategy for excitotoxicityrelated neurological disorders would selectively target only those NMDARs or their downstream signaling components associated with neuronal death, while sparing the cell survival-promoting NMDAR population.

## 4 NMDAR Signaling at the Postsynaptic Density

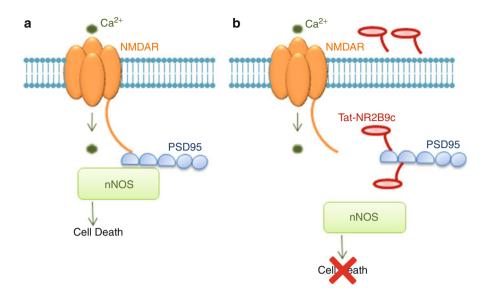
At the postsynaptic membrane of central excitatory synapses, ionotropic glutamate receptors are localized into specialized MPCs in an electron-dense region termed the postsynaptic density (PSD) (Kennedy 1997; Sheng 2001). NMDARs are abundantly expressed in association with the PSD (Moon et al. 1994), which is composed of both membranous and cytoplasmic proteins (Ziff 1997; Okabe 2007). Scaffolding proteins, which are major components of the MPC, associate with each other and other MPC proteins via highly specific and often unique protein-protein interactions. Such interactions govern cell-to-cell adhesion, regulation of receptor clustering, and modulation of receptor function (Feng and Zhang 2009; Verpelli et al. 2012). PSD95 (also known as SAP90), the first PSD scaffolding protein to be identified (Cho et al. 1992), is a member of a larger group of scaffolding proteins known as membrane-associated guanylate kinase (MAGUK) proteins (Gardoni 2008; Zheng et al. 2011). MAGUKs contain modular protein interaction domains, such as SH3 (Src homology 3), GK (guanylate kinase), and PDZ (Postsynaptic density-95/Discs large homolog/Zona occludens-1) domains (Sheng and Sala 2001; Gardoni 2008; Zheng et al. 2011).

Many surface receptors, such as the NMDAR, bind to MAGUK PDZ domains via their C-terminal PDZ-binding domains. For example, through its first two PDZ domains, PSD95 can bind NMDAR NR2 subunits at their C-terminus PDZ-binding motif (ESDV) (Kornau et al. 1995; Niethammer et al. 1996). PSD95 connects NMDARs to intracellular signaling proteins, such as neuronal nitric oxide synthase (nNOS). PSD95 also binds directly to nNOS via a PDZ-PDZ interaction that involves the second PDZ domain of PSD95 (Brenman et al. 1996; Craven and Bredt 1998; Christopherson et al. 1999; Tezuka et al. 1999). By physically bringing together the NMDAR and nNOS, PSD95 allows Ca<sup>2+</sup> that permeates through NMDARs to preferentially induce the activation of nNOS and thus couples NMDAR activity to the production of nitric oxide (NO), a signaling molecule that mediates NMDAR-dependent excitotoxicity (Dawson et al. 1991, 1993). Disrupting the NMDAR-PSD95-nNOS complex in cultured cortical neurons using antisense oligonucleotides against PSD95 selectively attenuated NMDAR-mediated excitotoxicity, without affecting NMDAR expression or function (Sattler et al. 1999).

## 5 NMDAR NR2B Signaling: Downstream Effectors of Excitotoxicity as Therapeutic Targets in Neurological Diseases

## 5.1 NR2B-PSD95-nNOS Signaling Complex

Because suppression of PSD95 using antisense oligonucleotides is impractical in the treatment of excitotoxicity-related disorders in humans, an alternative strategy



**Fig. 1** The NMDAR-PSD95-nNOS signaling complex. (a) By binding to both the NMDAR and nNOS, PSD95 brings nNOS in close proximity to the NMDAR and allows Ca<sup>2+</sup> entering the neuron through the NMDAR to preferentially induce the activation of nNOS, leading to cell death. (b) The Tat-NR2B9c peptide disrupts the NMDAR-PSD95-nNOS complex, dissociating NMDARs from downstream neurotoxic signaling, without blocking synaptic activity or calcium influx

for disrupting the NR2B-PSD95-nNOS signaling complex is to use drug compounds that block protein interactions within this complex. One such drug compound is NR2B9c (Aarts et al. 2002). The NR2B9c peptide, comprising the nine C-terminal residues of NR2B (KLSSIESDV), can be rendered cell permeant by fusion to the 11 residue protein transduction domain of the human immunodeficiency virus type 1 (HIV-1) Tat protein (Schwarze et al. 1999). This produces a 20-amino acid peptide termed Tat-NR2B9c. This peptide permeates through cell membranes and disrupts the intracellular interaction of NMDARs with PSD95 (See Fig. 1). This effectively dissociates NMDARs from downstream neurotoxic signaling. However, unlike NMDAR antagonists, this occurs without blocking synaptic activity or calcium influx. This approach is feasible in vivo: when applied either before, 1 or 3 h after an insult, Tat-NR2B9c protected cultured neurons from excitotoxicity and reduced transient middle cerebral artery occlusion (MCAO)-mediated ischemic brain damage in rats (Aarts et al. 2002; Sun et al. 2008).

A proteomic and biochemical analysis of the interactions of Tat-NR2B9c with most or all PDZs in the human genome revealed that the disruption of the NMDAR-PSD95-nNOS complex by Tat-NR2B9c was highly specific. Because the peptide targets the first and second PDZ domains of PSD95, where both the NMDAR and nNOS bind, Tat-NR2B9c inhibited the interaction between both PSD95 and NR2 subunits as well as between PSD95 and nNOS (Cui et al. 2007).

Importantly, Tat-NR2B9c administered 3 h after both permanent and transient focal ischemia reduced infarct volumes in rats and improved long-term neurobehavior, including sensorimotor function, emotionality, and cognition (Sun et al. 2008), raising the possibility for clinical usefulness in a wide therapeutic window. Further downstream, the consequences of administering Tat-NR2B9c are to ameliorate excitotoxic neuronal loss in vitro and ischemic cortical damage in vivo by impairing pro-death p38 signaling, without affecting NMDAR-mediated prosurvival pathways involving CREB or Akt (Soriano et al. 2008; Martel et al. 2009). Other studies have proposed that compounds other than Tat-NR2B9c that target the NMDAR-PSD95 interaction may also have neuroprotective effects. Disruption of the PSD95-nNOS interaction using a small molecule inhibitor (ZL006) is suggested to be neuroprotective in ischemia (Zhou et al. 2010). In addition, the use of a dimeric inhibitor, Tat-NPEG4(IETDV)<sub>2</sub> (Tat-*N*-dimer), which binds the tandem PDZ1-2 domain of PSD95, reduced infarct volume in mice subjected to cerebral ischemia (Bach et al. 2012).

Recently, Tat-NR2B9c was used for the treatment of stroke in cynomolgus macaques, higher-order gyrencephalic nonhuman primates, which bear anatomical and behavioral similarities to humans (Cook et al. 2012). Animals treated with Tat-NR2B9c up to 3 h after MCAO showed significant reduction in infarct volume, as measured by magnetic resonance imaging and histology. In addition, Tat-NR2B9c-treated macagues showed improved neurobehavioral outcomes assessed using the nonhuman primate stroke scale (NHPSS). This neuroprotection preserved cellular functionality, as gauged by the capacity for gene transcription in ischemic brain tissue. The 3-h time frame for neuroprotection with Tat-NR2B9c is significant as it suggests a broader clinical applicability. Tat-NR2B9c has already entered clinical trials; the ENACT (Evaluating Neuroprotection in Aneurysm Coiling Therapy) Phase 2 clinical trial (ClinicalTrials.gov number, NCT00728182) to determine the safety and efficacy of Tat-NR2B9c in reducing embolic strokes in patients undergoing endovascular repair of brain aneurysms was recently successfully completed (Hill et al. 2012). The encouraging results from this trial warrant further investigation of Tat-NR2B9c in the clinical treatment of stroke.

Since the first study (Aarts et al. 2002) utilizing blocking peptides against PSD95 in stroke, subsequent work using similar strategies has been carried out to investigate other excitotoxicity-mediated neurological diseases, such as epilepsy, AD, and neuropathic pain. Administration of Tat-NR2B9c in rats 3 h after the termination of status epilepticus reduced cell loss in regions of the hippocampus (Dykstra et al. 2009). Perturbing the NR2B-PSD95 association with Tat-NR2B9c ameliorated A $\beta$ -mediated toxicity in vitro and was sufficient to prevent lethality and memory deficits in an AD mouse model (Ittner et al. 2010). The importance of the NR2B9c-PSD95-nNOS signaling complex in neuropathic pain has been demonstrated using a variety of genetic, biochemical, and proteomic techniques. These include the use of antisense oligonucleotides against PSD95 (Tao et al. 2001, 2003), small molecule inhibitors that block the interaction between PSD95 and nNOS (Florio et al. 2009), or Tat-fusion peptides comprising regions of PSD95 (Tao et al. 2008), nNOS (Florio et al. 2009), or the NMDAR NR2B subunit (D'Mello et al. 2011).

## 5.2 NR2B-ND2-Src Signaling Complex

NMDAR activity is governed in part by the balance between phosphorylation and (Salter and Kalia 2004). dephosphorylation Tvrosine phosphorylation upregulates the activity of NMDARs (Wang and Salter 1994; Zheng et al. 1998), and the nonreceptor tyrosine kinase Src (Ohnishi et al. 2011) can mediate this upregulation. Within the NMDAR complex, Src, along with the phosphotyrosine phosphatase STEP (striatal enriched tyrosine phosphatase), act in opposition to phosphorylate or dephosphorylate the NMDAR, respectively. STEP<sub>61</sub>, the membrane-associated STEP isoform (Sharma et al. 1995; Bult et al. 1996), suppresses NMDAR activity by opposing the actions of Src (Pelkey et al. 2002). Src-mediated enhanced phosphorylation of NMDARs is associated with synaptic plasticity (Rostas et al. 1996; Salter 1998) as well as pathophysiological conditions such as ischemia (Cheung et al. 2003), HD (Song et al. 2003), epilepsy (Sanna et al. 2000; Moussa et al. 2001), and neuropathic pain (Salter and Pitcher 2012). Inhibition of Src activity may attenuate brain injury in ischemia (Jiang et al. 2008; Liang et al. 2009), supporting a role for Src in excitotoxic processes.

Src is anchored to the NMDAR via the adaptor protein NADH dehydrogenase 2 (ND2) (Gingrich et al. 2004). ND2 brings Src in close proximity to the NMDAR and allows Src to upregulate NMDAR function through phosphorylation of NR2 subunits. Disruption of the NMDAR-ND2-Src signaling complex has been shown to suppress pain behaviors. A 10-amino acid peptide, Src40-49, comprising the region of Src needed to bind ND2, was able to inhibit the binding of Src to ND2, thus releasing Src from the NMDAR complex and inhibiting Src-mediated enhancement of NMDAR activity (Liu et al. 2008). Importantly, uncoupling of Src from the NMDAR complex with the membrane-permeable Src40-49Tat fusion peptide suppressed pain hypersensitivity induced by inflammation and peripheral nerve injury (see Fig. 2). Thus, by regulating the function of NMDARs through phosphorylation and dephosphorylation, Src and STEP play vital roles in both physiological as well as pathophysiological glutamatergic neurotransmission. Furthermore, both Src and STEP themselves are subject to regulation. For example, differential regulation of  $STEP_{61}$  may play an important role in mediating the dual roles of the NMDAR in cell survival and death (Xu et al. 2009). Synaptic NMDAR activation promotes the ubiquitination and degradation of STEP<sub>61</sub>, concomitant with ERK1/2 activation and cell survival. In contrast, extrasynaptic NMDAR stimulation leads to calpainmediated proteolysis of STEP<sub>61</sub>, activation of p38, and cell death. Aberrant regulation of both Src and STEP may therefore be involved in NMDAR-mediated pathophysiological conditions (Hossain et al. 2012; Salter and Pitcher 2012). Although targeting STEP or the Src:ND2 interaction has not been tested in ischemia, it is another example of a protein-protein interaction that may be targeted therapeutically in disorders that depend on NMDAR overactivity.

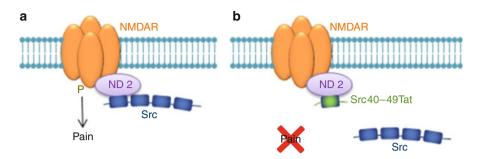


Fig. 2 The NMDAR-ND2-Src signaling complex. (a) Src is anchored within the NMDAR complex via the adaptor protein NADH dehydrogenase 2 (ND2). By bringing Src in close proximity to the NMDAR, ND2 allows Src to phosphorylate the receptor and enhance NMDAR activity, resulting in hypersensitivity to pain. (b) Using a Src40–49Tat peptide to block the interaction between Src and ND2 dissociates Src from ND2, thereby inhibiting Src-mediated upregulation of NMDAR activity and suppressing pain behaviors (Figure adapted from Liu et al. 2008)

## 5.3 Other Signaling Pathways Implicated in NMDAR-Mediated Neurotoxicity

Other signaling pathways downstream of NMDARs have been identified to play roles in excitotoxic signaling. For example, during cerebral ischemia, deathassociated protein kinase 1 (DAPK1) is recruited to extrasynaptic NMDAR NR2B subunits in the cortex of mice (Tu et al. 2010). DAPK1 directly binds to NR2B, phosphorylates it, and enhances NMDAR channel conductance, leading to increased excitotoxic signaling. Peptides that inhibit the interaction between DAPK1 and the NMDAR NR2B subunit reduced brain infarction and improved neurological function in mice. DAPK1 itself is subject to regulation by other signaling molecules, such as ERK1/2, which may affect NMDAR activity (Liu et al. 2012). Another signaling protein involved in cell death downstream from the NMDAR is PTEN (phosphatase and tensin homolog deleted on chromosome 10). PTEN is recruited selectively to NMDAR NR2B subunits (Ning et al. 2004), potentiates neurotoxic NMDAR activity, and inhibits the PI3K (phosphatidylinositol 3-kinase) survival pathway, thus contributing to excitotoxic neuronal death in stroke. PTEN-induced kinase 1 (PINK1) in a ubiquitous kinase thought to function in cell survival-promoting pathways (Wilhelmus et al. 2012). Overactivation of NR2B-containing NMDARs may contribute to ischemic cell death by suppressing PINK1-dependent survival signaling (Shan et al. 2009). Thus, drug compounds that are able to increase PINK1 levels and activity may prove to be neuroprotective in NMDAR-mediated neurotoxicity.

Activation of NMDARs has been linked to the modulation of several transcription factors that can mediate either pro-survival or pro-death pathways (Hardingham et al. 2002; Zou and Crews 2006; Lai et al. 2011), and these may be useful targets for therapeutic intervention. One example is the transcription factor

SREBP-1 (sterol regulatory element-binding protein-1) (Taghibiglou et al. 2009), which typically controls lipid biosynthesis genes but whose activation is essential in NMDAR-mediated excitotoxic neuronal death in both in vitro and in vivo models of stroke. Therapies aimed at pro-death transcription factors downstream of NMDAR-mediated excitotoxic pathways may offer some novel therapeutic targets in the treatment of neurological diseases. Since activation of these transcription factors is often delayed, relative to NMDAR activation, neuroprotective strategies targeting these proteins may offer a wider therapeutic window if such pathways participate in the cell injury process.

Cell death pathways also exist that may occur in parallel with and converge upon NMDAR-mediated excitotoxic signaling mechanisms (Tymianski 2011). TRPM7 and TRPM2 are members of the transient receptor potential (TRP) channel superfamily (Nilius and Owsianik 2011) that are widely expressed and function in diverse cellular processes, including death, survival, and proliferation. Both TRPM2 and TRPM7 are nonselective cation channels that are activated by oxidative stress and are thought to play significant roles in ischemic cell death (Aarts et al. 2003a; MacDonald et al. 2006; Sun et al. 2009; Xie et al. 2010). TRPM2 and TRPM7 may represent additional Ca<sup>2+</sup> influx pathways, other than NMDARs, in mediating neurotoxicity in ischemia. Neurotoxic signals such as reactive oxygen species downstream of NMDARs may in turn activate TRPM2 and TRPM7 channels to form a positive feedback loop that perpetuates ischemic cell death.

# 6 Conclusion

NMDARs have important physiological roles mediating synaptic plasticity and neurodevelopment. However, overactivation of NMDARs leads to excitotoxicity, common to many neurological diseases such as ischemic stroke, epilepsy, neuropathic pain, traumatic brain and spinal injuries; ocular disorders such as glaucoma; as well as neurodegenerative diseases, including amyotrophic lateral sclerosis, Parkinson's, Huntington's, and Alzheimer's diseases. Antagonists that directly inhibit NMDARs have limited use clinically, possibly due to intolerable side effects at anti-excitotoxic doses. Intracellular signaling pathways that couple to NMDAR activation can promote either cell survival or cell death, depending on NMDAR subunit composition and/or subcellular localization. Extrasynaptic NMDARs, and especially those receptors containing NR2B subunits, engage in cell death pathways upon excessive glutamate stimulation. Conversely, activation of synaptically localized NMDARs, and especially those receptors containing NR2A subunits, may result in neuroprotection. Therapeutic strategies targeting NMDAR-mediated neurotoxicity should preferentially block pro-death signaling while sparing pro-survival signals. Since neuroprotectants that selectively target the molecular components involved in pro-death signaling downstream of the NMDAR do not directly inhibit NMDARs themselves, they may be better tolerated than traditional NMDAR antagonists.

The last decade of scientific research has shed light on the molecular mechanisms involved in NMDAR-mediated excitotoxicity and has revealed additional targets for neuroprotective strategies in the treatment of neurological disorders; however, many questions remain unanswered. First, in addition to NR1/NR2A and NR1/NR2B diheteromeric NMDARs, triheteromeric receptors comprising NR1/NR2A/N2B are also believed to exist. The roles of these triheteromeric receptors, in both health and disease, have yet to be elucidated. Second, several signaling molecules downstream of NMDARs, such as nNOS, DAPK1, and PTEN, play roles in excitotoxic signaling to promote cell damage and death. Whether and how these signaling molecules cross talk with one another is not known. Third, in conditions such as ischemia, excessive  $Ca^{2+}$  influx through NMDARs leads to the formation of reactive oxygen species, which can activate other plasma membrane channels, such as TRPM2 and TRPM7 channels; these channels thus represent additional therapeutic targets. Lastly, a major part of our understanding about NMDAR-mediated cell death pathways comes from models of cerebral ischemia. Future work will need to determine whether these pathways also play a role in other neurological diseases that involve excitotoxic mechanisms and whether preclinical neuroprotective therapies can be translated to clinical use.

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# Neuroprotection by Kynurenine Metabolites

# Zsófia Majláth, Levente Szalárdy, Dénes Zádori, Péter Klivényi, Ferenc Fülöp, József Toldi, and László Vécsei

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#### Abstract

The overexcitation of excitatory amino acid receptors generates excitotoxicity, which plays an important role in the pathophysiology of various neurodegenerative disorders such as Parkinson's and Alzheimer's diseases, stroke, and epilepsy. The prevention of excitotoxic neuronal damage is therefore a major

e-mail: vecsei.laszlo@med.u-szeged.hu

Z. Majláth • L. Szalárdy • D. Zádori • P. Klivényi

Department of Neurology, University of Szeged, Szeged, Hungary

F. Fülöp

Institute of Pharmaceutical Chemistry and Research Group of Stereochemistry, Hungarian Academy of Sciences, University of Szeged, Szeged, Hungary

J. Toldi

Department of Physiology, Anatomy and Neuroscience, University of Szeged, Szeged, Hungary

Neuroscience Research Group of the Hungarian Academy of Sciences and University of Szeged, Szeged, Hungary

L. Vécsei (⊠) Department of Neurology, University of Szeged, Szeged, Hungary

Neuroscience Research Group of the Hungarian Academy of Sciences and University of Szeged, Szeged, Hungary

objective in neuroprotective strategies. The kynurenine pathway, the main route of the tryptophan metabolism, produces both neuroprotective and neurotoxic metabolites. Kynurenic acid is a broad-spectrum endogenous antagonist of ionotropic excitatory amino acid receptors, and hence it can prevent excitotoxic neuronal damage. The kynurenine metabolites are additionally involved in the regulation of glutamatergic and cholinergic neurotransmission. The pathogeneses of several neurodegenerative disorders have been shown to involve multiple imbalances in the kynurenine metabolism. Kynurenine pathway metabolites may provide important targets for future neuroprotective therapeutic strategies.

#### Keywords

Glutamate excitotoxicity • Kynurenine pathway • Neurodegeneration • Neuroprotection

#### List of Abbreviations

3-HANA	3-hydroxyanthranilic acid
AMPA	alpha-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid
ANA	anthranilic acid
3-HK	3-hydroxykynurenine
EAA	excitatory amino acid
IDO	indoleamine 2,3-dioxygenase
KAT	kynurenine aminotransferase
KMO	kynurenine 3-monooxygenase
KYNA	kynurenic acid
L-KYN	L-kynurenine
NAD	nicotinamide adenine dinucleotide
NMDA	N-methyl-D-aspartate
QUIN	quinolinic acid
TDO	tryptophan 2,3-dioxygenase

# 1 Introduction

The term excitotoxicity refers to pathological processes in which excitatory amino acid (EAA) receptors are overactivated, thereby initiating further pathological processes and ending with the death of neurones. This mechanism is involved in the pathophysiology of diverse neurological diseases such as stroke or epilepsy, traumatic brain injury, and various neurodegenerative disorders including Parkinson's disease and Huntington's disease. One of the most important substances participating in these processes is glutamate, which is also the most important excitatory neurotransmitter in the mammalian brain. The results of one of the first studies implicating the toxic effect of glutamate were published in 1957, when Lucas and Newhouse observed that the subcutaneous injection of monosodium glutamate destroyed the neurones in the inner layers of the retina in newborn mice (Lucas and Newhouse 1957). It was later

established that this phenomenon occurs not only in the retina but in several brain regions too: it is restricted to postsynaptic neurones, and glutamate agonists may induce the same effect (Olney 1969). These findings led to emergence of the term excitotoxicity. Glutamate acts on different types of ionotropic and G protein-coupled (metabotropic) receptors. These receptors play essential roles in the normal functioning of the brain, but their excessive stimulation may trigger an extensive influx of calcium (Ca<sup>2+</sup>) to the neurones, setting into motion various pathological processes, which lead to damage to the cell structures and terminate in cell death (Arundine and Tymianski 2003; Krieglstein 1997). Excitotoxicity is known to be involved in the pathomechanism of a number of neurological diseases, and the search for neuroprotective agents which may prevent neuronal damage has long been an important aim in neuroscience. The kynurenine pathway, the main route in the tryptophan catabolism, involves both an EAA receptor antagonist and an agonist (Beadle et al. 1947), and kynurenines are involved in various physiological and pathophysiological processes, therefore standing at the focus of research interest (Schwarcz 2004).

#### 2 Glutamatergic Excitotoxicity

Ionotropic receptors activated by glutamate can be divided into three major types identified by their preferred agonists: N-methyl-D-aspartate (NMDA), kainate, and alpha-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA), each of them with several subtypes. All three families of ionotropic glutamate receptors allow the permeation of sodium (Na<sup>+</sup>) and potassium (K<sup>+</sup>), whereas the NMDA-type and some AMPA-type channels also display Ca<sup>2+</sup> permeability. Metabotropic receptors activated by glutamate exert their effects through the activation of second messenger systems also leading to a cytosolic Ca<sup>2+</sup> overload. The different receptor families contribute to the excitotoxic processes to different extents, and there is evidence that the NMDA receptors play a pivotal role. AMPA receptors have a heterotetrameric structure built up from four homologous subunits (GluR1-4) (Hollmann and Heinemann 1994). All of the subunits consist of a large extracellular N-terminal domain followed by four transmembrane domains ending in an intracellular C-terminal segment. The second transmembrane domain is responsible for the formation of the cation-permeable channel (Ozawa et al. 1998; Madden 2002). The glutamate-binding domain is formed by a part of the N-terminal domain together with the third membrane domain (Hollmann et al. 1994). The C-terminal domain is responsible for the interaction with the intracellular proteins and contains phosphorylation sites too (Carvalho et al. 2000). The subunits may exist in two distinct forms, flip and flop, which are formed through alternative splicing (Sommer et al. 1990). The activation of AMPA and kainate receptors allows the entry of Na<sup>+</sup> to the neurones, triggering rapid excitatory neurotransmission in the brain. Additionally in the hippocampus, cortex, and retina, various neurones express AMPA receptors which are also permeable to Ca<sup>2+</sup>. Under physiological conditions, the number of Ca<sup>2+</sup>-permeable AMPA receptors present in the hippocampal pyramidal neurones (HPNs) is relatively low, but in consequence of ischaemia, this number undergoes a significant increase (Pellegrini-Giampietro et al. 1997).

NMDA receptors are heteromeric complexes consisting of subunits of three major types: the NR1, NR2, and NR3 families. In mammalian tissues, functional NMDA receptors are expressed only when they contain at least one NR1 and one NR2 subunit (Moriyoshi et al. 1991; Ciabarra et al. 1995). NMDA receptors are slow-gating ion channels that are highly permeable to Ca<sup>2+</sup>. The C-terminal domains of NMDA receptor subunits have an important role in NMDA receptor interactions with intracellular synaptic and cytoskeletal proteins, and these protein-protein interactions initiate various downstream signalling pathways (reviewed by Arundine and Tymianski 2004). It has been demonstrated experimentally that glutamate excitotoxicity has two components: immediate cell damage which is Na<sup>+</sup>- and Cl<sup>-</sup>-dependent and a delayed neuronal degeneration process which can be related to the presence of Ca<sup>2+</sup>. Na<sup>+</sup> influx causes neuronal damage even in the absence of  $Ca^{2+}$ , through the generation of toxic swelling in hippocampal and retinal cells (Rothman 1985; Olney et al. 1986). The neuronal swelling can be avoided by the removal of extracellular Na<sup>+</sup>, but neurones exposed to glutamate will still develop neurodegenerative processes unless the extracellular Ca2+ is also removed (Berdichevsky et al. 1983; Choi et al. 1987). Other possible ways for Ca<sup>2+</sup> to enter the cells are voltage-dependent  $Ca^{2+}$  channels and  $Na^+/Ca^{2+}$  exchangers. Release from internal pools may contribute to toxic Ca<sup>2+</sup> concentrations. This can occur through mitochondrial damage or dysfunction. An increased intracellular Ca<sup>2+</sup> level initiates pathological processes such as proteolysis, protein phosphorylation, lipolysis, the activation of nitric oxide synthases, and the generation of free radicals which may destroy neurones (Szydlowska and Tymianski 2010). The activation of proteases, lipases, and endonucleases results in damage to cell structures, degradation of the cytoskeleton and DNA and a mitochondrial dysfunction, and finally leads to cell death.

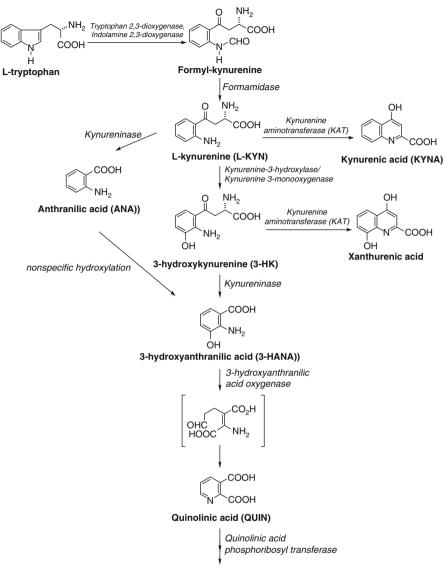
If neurones suffer an energy impairment, a partial membrane depolarization occurs, which leads to the removal of  $Mg^{2+}$  blocking the NMDA receptors, allowing glutamate to exert its downstream effects under physiological concentrations, contributing to neurodegenerative processes (Novelli et al. 1988).

As excitotoxic mechanisms are involved in pathological neurone loss, EAA receptor antagonism has long been at the focus of interest for neuroscientists searching for neuroprotective agents. The early experiments with NMDA antagonists revealed that they are able to prevent the neurotoxic effects of NMDA (Olney et al. 1979) and ibotenate (Schwarcz et al. 1982). In animal models, NMDA antagonism proved to have anticonvulsive properties (Croucher et al. 1982) and to be able to prevent neuronal damage in cerebral ischaemia (Simon et al. 1984). Subsequent to these convincing results, numerous experiments have demonstrated the neuroprotective efficacy of NMDA antagonism.

#### 3 The Kynurenine Pathway and Its Metabolites

#### 3.1 The Kynurenine Pathway

L-Tryptophan, one of the ten essential amino acids, serves as a precursor of several biologically active molecules. Besides its involvement in protein synthesis, it can be



Nicotinamide adenine dinucleoid (NAD)



metabolized in different pathways. The serotonin pathway is responsible for the synthesis of the important neurotransmitter serotonin, which can be converted to melatonin. In the human brain, as in most mammalian tissues, 95 % of tryptophan is catabolized through the kynurenine pathway (reviewed by Vamos et al. 2009) (Fig. 1). This metabolic cascade is responsible for the synthesis of nicotinamide adenine dinucleotide (NAD) and NAD phosphate (NADP), coenzymes which are essential for many

biological processes, such as the control of the energy metabolism of cells, DNA repair, and transcription. The other metabolites of this pathway, collectively termed "kynurenines," have been shown to play important roles in the nervous and immune systems and to be involved in the pathophysiology of a growing number of neurological diseases (Ogawa et al. 1992; Stoy et al. 2005; Hartai et al. 2005; Robotka et al. 2008b; Guidetti et al. 2006).

The first step in this cascade is the enzymatic degradation of tryptophan by indoleamine 2,3-dioxygenase (IDO) or tryptophan 2,3-dioxygenase (TDO) to formylkynurenine, which is converted by arylformamidase to L-kynurenine (L-KYN). The pathway then divides into three main branches. The first branch ends with the synthesis of kynurenic acid (KYNA) through the transamination of L-KYN, a metabolic step catalyzed by the kynurenine aminotransferases (KATs) (Rossi et al. 2008). It was first believed that there are two main KAT isoforms in the brain (Okuno et al. 1991). The KATs are mainly localized in the astrocytes (Du et al. 1992), but they are also present in neurones in the hippocampus, in the striatum (Knyihar-Csillik et al. 1999), and in most of the neurones in the medulla and spinal cord (Kapoor et al. 1997). Early studies indicated that under physiological pH conditions, KAT-II has higher substrate specificity for KYN than KAT-I and that in most brain regions, KYNA can be attributed primarily to KAT-II activity (Okuno et al. 1991; Guidetti et al. 1997; Kiss et al. 2003). The results of recent experiments relating to the pH profile of human KAT-I indicated that under physiological pH conditions, it is able to catalyze L-KYN to KYNA (Han et al. 2004); hence, it may also play an important role in kynurenine metabolism. Recently, another member of the KAT family was discovered, the KAT-III isoform, which exhibits a number of similarities to KAT-I but appears to have no activity in the human brain (Yu et al. 2006). Whereas the main enzyme of KYNA synthesis in the human and the rat brain is KAT-II, recent studies revealed that the most important enzyme in mice is KAT-IV, which has proved to be identical to mitochondrial aspartate aminotransferase (mtAAT) (Guidetti et al. 2007; Han et al. 2010). The four KAT isoforms have different pH optima and temperature preferences (Han et al. 2010).

L-KYN can also serve as a substrate for kynureninase, giving rise to anthranilic acid (ANA). The third main branch of the kynurenine pathway comprises a sequence of enzymatic steps which ends in the synthesis of NAD. This cascade is responsible for the formation of free radical generators 3-hydroxykynurenine (3-HK) and 3-hydroxyanthranilic acid (3-HANA) and the neurotoxic compound quinolinic acid (QUIN). The first of these steps is to yield 3-HK that involves the action of kynurenine 3-monooxygenase (KMO; kynurenine-3-hydroxylase), which is then degraded by kynureninase into 3-HANA. This metabolite, which may also be generated from ANA, is then converted by 3-HANA-oxygenase to 2-amino-3-carboxymuconate-semialdehyde, an unstable intermediate which is transformed nonenzymatically into QUIN. Finally, QUIN is transformed in consecutive steps involving the enzymatic action of phosphoribosyltransferase to furnish nicotinamide and NAD.

The enzymatic machinery for this pathway is present in both the brain and the periphery, but the metabolic activity differs under physiological and pathophysiological conditions. Under physiological conditions, this pathway has a much higher capacity at the periphery, while during inflammatory processes localized to the central nervous system (CNS), more than 98 % of the L-KYN and QUIN result from local synthesis in the brain (Kita et al. 2002). KYNA and QUIN are present in nanomolar concentrations in the human cerebrospinal fluid (CSF) (Moroni 1999; Schwarcz and Pellicciari 2002; Erhardt et al. 2004), while the concentration of KYNA in the human brain tissue is assumed to be 0.2–1.5  $\mu$ M (Moroni et al. 1988; Turski et al. 1988). The neuroactive metabolites KYNA and QUIN cross the blood–brain barrier to only a limited extent, whereas their precursor L-KYN is able to cross it well (Fukui et al. 1991).

Within the CNS, the enzymes are differently distributed between the different subtypes of glial cells. Many studies have indicated that the astrocytes harbor hardly any KMO and therefore contribute more significantly to the synthesis of KYNA. In contrast, the microglial cells contain little KAT and hence are primarily responsible for the formation of QUIN (Guillemin et al. 2001; Lehrmann et al. 2001; Kiss et al. 2003).

#### 3.2 Quinolinic Acid and Other Kynurenines with Neurotoxic Properties

QUIN is a weak competitive agonist of the NMDA receptors containing NR2A and NR2B subunits (Stone and Perkins 1981; de Carvalho et al. 1996). OUIN has been reported to be involved in the pathogenesis of several neurodegenerative disorders (Schwarcz et al. 1984; Klivenyi et al. 2004). Early studies yielded evidence that the administration of QUIN leads to a neurotoxic effect and may result in neuronal death in cortical cell cultures (Kim and Choi 1987). In vivo it has been observed that QUIN produces axon-sparing lesions in the rat brain (Schwarcz et al. 1983), which are similar to those seen in Huntington's disease (Beal et al. 1986). Intrastriatal QUIN injection is therefore a widely used experimental animal model for the study of Huntington's disease. It has proved to be neurotoxic in a number of studies conducted in recent years, but its neurotoxic effect involves several processes besides NMDA receptor agonism. It has become clear from the results that QUIN causes endogenous antioxidant depletion, contributes to the generation of free radicals (Behan et al. 1999; Rodriguez-Martinez et al. 2000), and also induces lipid peroxidation (Rios and Santamaria 1991). Moreover, QUIN has been confirmed to influence the glutamate metabolism, as it can enhance presynaptic glutamate release and inhibit glutamate uptake by astrocytes (Connick and Stone 1988; Tavares et al. 2002). Plasma levels of QUIN are elevated in Parkinson's disease patients (Ravikumar et al. 2000). Human studies revealed that QUIN levels are also increased in early stages of Huntington's disease (Guidetti et al. 2004).

3-HK and 3-HANA are potent neurotoxic agents experimentally demonstrated to be able to induce apoptosis in cultured cells (Morita et al. 1999). 3-HK and 3-HANA also contribute to the production of free radicals and elevation of the oxidative stress level. Under physiological conditions, the concentration of 3-HK is

within the nanomolar range, but there is evidence that under certain pathological conditions, it can reach micromolar levels (Okuda et al. 1998; Pearson and Reynolds 1991). The level of 3-HANA, a metabolite produced from 3-HK or ANA, is elevated in Huntington's disease and Parkinson's disease (Ogawa et al. 1992; Pearson and Reynolds 1992).

#### 3.3 The Endogenous Neuroprotectant Kynurenic Acid

KYNA, one of the neuroactive metabolites of the kynurenine pathway, displays broadspectrum antagonistic properties of ionotropic EAA receptors (Stone and Connick 1985). KYNA is an endogenous glutamate receptor antagonist of AMPA, NMDA, and kainate types, with the highest affinity for NMDA receptors (Perkins and Stone 1985; Swartz et al. 1990). At low micromolar concentrations (EC50 =  $7.9-15 \mu$ M), it has a high affinity for the strychnine-insensitive glycine-binding site of the NMDA receptors and blocks it in a competitive way, while at 10–20 times higher concentrations, it exhibits an antagonistic effect of the glutamate-binding site too (EC50 =  $200-500 \mu$ M) (Birch et al. 1988; Kessler et al. 1989). High concentrations of KYNA exert an anticonvulsant effect and protect the neurones from excitotoxic injury.

In vitro studies have revealed that KYNA may be capable of the direct activation of the G protein-coupled receptor GPR35 (Wang et al. 2006; Ohshiro et al. 2008). KYNA was earlier recognized to display antagonistic properties at the AMPA receptors, and it has only recently become clear that, depending on its concentration, it may also have another mechanism of action. In micromolar concentrations, KYNA inhibits AMPA receptors, whereas in nanomolar concentrations, it acts as a facilitator, binding to a positive modulatory-binding site at the AMPA receptor (Prescott et al. 2006; Rozsa et al. 2008). This dose-dependent dual effect indicates that KYNA may play an important role in the regulation of neurotransmission. KYNA has proved to exert an antagonistic effect on the presynaptic  $\alpha$ 7-nicotinic acetylcholine ( $\alpha$ 7-nACh) receptor too, inhibiting its activity in a noncompetitive manner, while increasing the expression of non- $\alpha$ 7-nACh receptors (Hilmas et al. 2001). This effect may contribute to the generation of the sensory deficit caused by KYNA (Shepard et al. 2003; Chess and Bucci 2006), and the evidence suggests that it is mediated by its binding to sites located on the N-terminal domain of the  $\alpha$ 7-nACh receptor subunit (Pereira et al. 2002). Recent studies have revealed that KYNA in low concentrations (30-100 nM) inhibits glutamate release at the presynaptic nACh receptors, thereby contributing to a neuroprotective effect (Carpenedo et al. 2001; reviewed by Zadori et al. 2011). These results support the assumption that the kynurenine pathway is involved in the regulation of glutamatergic and cholinergic neurotransmission in the brain.

An increased level of KYNA was earlier proved to have sedative and anticonvulsive effects (Carpenedo et al. 1994). It was later demonstrated that it is able to reduce neuronal damage in brain ischaemia (Cozzi et al. 1999). There is a growing body of evidence that alterations in the kynurenine metabolism play important roles in the pathomechanisms of neurodegenerative disorders. Human postmortem studies have revealed elevated 3-HK levels in the substantia nigra and the putamen of Parkinson's disease patients, while the ratio L-KYN/3-HK is decreased (Ogawa et al. 1992). In Huntington's disease patients, the KYNA levels are significantly decreased in the striatum, cortex, and CSF (Beal et al. 1990, 1992). KYNA levels are elevated in Alzheimer's disease (Baran et al. 1999) and in schizophrenia (Schwarcz et al. 2001; Nilsson et al. 2005), playing a part in the cognitive decline. A number of experiments have demonstrated an elevated KYNA level or L-KYN exerts a neuroprotective effect in brain ischaemia (Nozaki and Beal 1992; Gigler et al. 2007; Robotka et al. 2008a; Sas et al. 2008).

#### 4 Conclusion

Excitotoxicity is a major factor in the pathomechanism of several neurological disorders. The overstimulation of EAA receptors results in an excessive influx of  $Ca^{2+}$  into the cells, subsequently initiating downstream pathological processes, resulting in damage to the cell structure, mitochondrial dysfunction, and an energy impairment leading to apoptosis. The kynurenine pathway is the main route of the metabolism of tryptophan in the mammalian brain. This pathway generates various neuroactive compounds which play an important role in the regulation of neurotransmission. There is accumulating evidence that imbalances in the kynurenine metabolism are involved in the pathomechanism of several neurodegenerative disorders, thereby providing a valuable target for future therapeutic strategies.

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# Neuroprotective Strategies in Amyotrophic Lateral Sclerosis: Modulation of Neurotransmittory and Neurotrophic Input to Motor Neurons

Pierre-François Pradat and Luc Dupuis

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#### Abstract

Amyotrophic lateral sclerosis (ALS) is an adult-onset degenerative disease characterized by the selective progressive death of lower and upper motor neurons. Although its primary cause remains unknown, multiple pathogenic pathways have been identified. The major pathological event in ALS is the degeneration of lower motor neurons (LMNs), and it is thought to be the ultimate cause of death. Thus, neuroprotection of LMNs is assumed to be a reasonable target for

P.-F. Pradat (🖂)

L. Dupuis

Département des Maladies du Système Nerveux, AP-HP, Groupe Hospitalier Pitié-Salpêtrière, UMR-678, INSERM-UPMC, Paris, France e-mail: pierre-francois.pradat@psl.aphp.fr

INSERM, U1118, Faculté de Médecine, Mécanismes centraux et périphériques de la neurodégénérescence, Université de Strasbourg, Strasbourg, France e-mail: ldupuis@unistra.fr

treating ALS. This assumption guided most of research effort in the last 20 years to develop neuroprotective strategies able to preserve the remaining LMNs. Three major possibilities have been explored and are presented in this review. First, it is thought that LMN survival and degeneration are regulated by neurotransmittory inputs that LMN integrates. Several experimental and imaging studies have shown an impairment of multiple neuronal types innervating motor neurons, in particular glycinergic interneurons, glutamatergic input from proprioceptive type Ia fibers, and serotonergic neurons. The abnormalities in glutamatergic input proved especially fruitful since they led to the discovery of riluzole. A second potential neuroprotective strategy would be to modulate neurotrophic input to motor neurons. Indeed, studies of motor neuron development have shown that their survival is governed by neurotrophic input from its muscle target or from surrounding glia. Thus, modulating neurotrophic support to LMN might be a therapeutic strategy, although the multiple clinical trials based on this possibility were up to now unsuccessful. A last possibility to provide neuroprotection in ALS would be to indirectly modulate the motor neuron survival through action on events occurring outside this cell type, in particular inflammation or abnormal energy metabolism.

#### Keywords

ALS • GABA • Glutamate • Glycin • Interneuron • Motor neuron • MRI • Neurotrophic factor • PET • Serotonin

#### List of Abbreviations

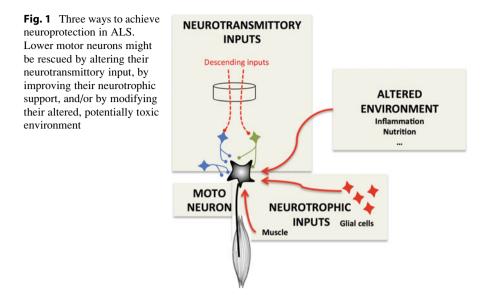
<sup>1</sup> H-MRS	Primary motor area, M1, proton magnetic resonance spectroscopy
5-HT	5-hydroxytryptamine
5-HTP	5-hydroxytryptophan
ALS	Amyotrophic lateral sclerosis
AMPA	α-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid
BDNF	Brain-derived neurotrophic factor
CNTF	Ciliary-derived neurotrophic factor
CSF	Cerebrospinal fluid
DTI	Diffusion tensor imaging
EAAT2	Excitatory amino acid transporter type 2
Exc.	Excitatory
GABA	Gamma-aminobutyric acid
GDNF	Glial cell line-derived neurotrophic factor
Glu	Glutamate
Gly	Glycine
IGF-1	Insulin-like growth factor-1
IN	Interneuron
inh.	Inhibitory
LMN	Lower motor neuron
mRNA	Messenger ribonucleic acid

MT	Magnetization transfer
NMDA	N-methyl-D-aspartate
PET	Positron emission tomography
PM	Premotor
SICI	Short-interval intracortical inhibition
SMA	Supplementary motor area
SOD1	Superoxide dismutase type 1
UMN	Upper motor neuron
VEGF	Vascular endothelial growth factor
VGLUT2	Vesicular glutamate transporter type 2

### 1 Introduction

Amyotrophic lateral sclerosis (ALS) is the major adult-onset motor neuron disease with an incidence of 2–3/100,000. ALS typically affects lower motor neurons (LMNs), which degenerate, leading to a progressively lethal paralysis. ALS symptoms might start in limbs (spinal onset, 2/3 of cases) or in muscles controlling speech and/or swallowing (bulbar onset, 1/3 of cases). Apart from LMN involvement, ALS is associated with degeneration of upper motor neurons (UMNs), and most patients develop UMN signs, in particular hyperreflexia and spasticity. A number of patients also display focal or more widespread frontal lobe disease (Schreiber et al. 2005). Other non-motor symptoms include major defects in energy metabolism, typified by weight loss, muscle wasting, and hypermetabolism (Dupuis et al. 2011). Death of ALS patients typically occurs within 2–5 years after diagnosis, but a huge heterogeneity among patients exists, with survival after diagnosis ranging from few weeks to more than 30 years (Chio et al. 2009).

The major pathological event in ALS being degeneration of LMNs, it is thought to be the ultimate cause of death, thus neuroprotection of LMNs is assumed to be a reasonable target for treating ALS. This assumption indeed guided most of research effort in the last 20 years to develop neuroprotective strategies able to preserve the remaining LMNs. Three major possibilities have been explored and will be developed sequentially in this chapter (Fig. 1). First, it is thought that LMN survival and degeneration is regulated by neurotransmittory inputs that LMN integrates, in particular glutamatergic input. This possibility proved especially fruitful since it culminated in the discovery of riluzole. Second, studies of LMN development have shown that LMN survival is governed by neurotrophic input from its muscle target or from surrounding glia. Thus, modulating neurotrophic support to LMN might be a therapeutic strategy. Third, most recent results in ALS field suggested that neuroprotection of LMN might be indirectly achieved through modulation of LMN independent events such as inflammation or nutrition.



#### 2 Neuroprotection Through Modulation of Neurotransmittory Inputs to Motor Neurons

In this first section, we will review how input to motor neurons might be modulated to achieve neuroprotection in ALS.

# 2.1 Major Inputs to Motor Neurons

Motor neuron activity is controlled by two types of inputs. First, activation of ionotropic receptors, either excitatory or inhibitory, will generate postsynaptic currents that summate at the soma. If the net resulting current is sufficient to elicit an action potential, then the motoneuron discharge, which triggers the muscle contraction. Second, neuromodulatory inputs, through metabotropic receptors, activate intracellular signalling pathways that modulate the responses to ionotropic receptors. Since electrical activity is critical in the survival of any neuron, it is likely that identifying the abnormalities of inputs to motor neurons and correct these through pharmacological means will provide therapeutic strategies.

Motor neurons receive inputs from the periphery (in particular proprioceptive neurons) and from central projections. A schematic drawing of inputs to LMNs is presented in Fig. 2. There are about 100,000 synaptic boutons per lumbar motor neurons in cat (Rekling et al. 2000). A majority of these boutons are inhibitory, either GABAergic or glycinergic. Such inhibitory connections come from Renshaw cells that mediate recurrent inhibition of motor neurons and from other spinal inhibitory interneurons, which integrate numerous descending and peripheral

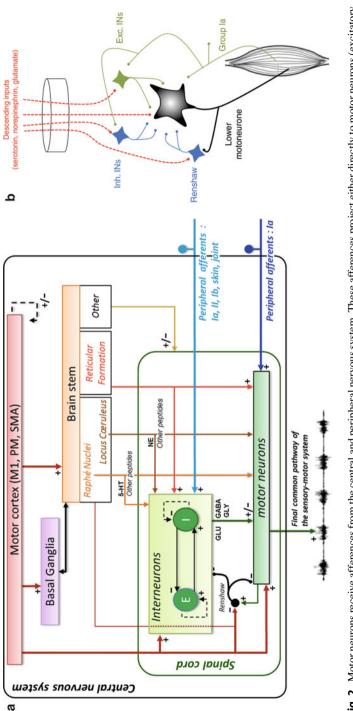


Fig. 2 Motor neurons receive afferences from the central and peripheral nervous system. These afferences project either directly to motor neurons (excitatory glutamatergic inputs) or indirectly through spinal intemeurons. Inputs from interneurons are mainly inhibitory, either either GABAergic or glycinergic, or excitatory, glutamatergic. Inhibitory interneurons include the Renshaw cells, glycinergic, which are activated by upper motor neurons and the interneurons, either GABAergic or glycinergic, which are activated by peripheral sensory neurons from skin and muscle spindles. Type Ia fibers coming from muscle spindles are glutamatergic and directly innervate motor neurons.

inputs. Inhibitory projections from brainstem nuclei and nucleus tractus solitarius also innervate spinal motor neurons.

About one fifth of synaptic boutons are glutamatergic, and include projections from type Ia muscle spindles and rubrospinal neurons. In rodent, corticospinal neurons do not directly project to motor neurons, but to spinal cord interneurons, thus only indirectly modulating motor neuron activity. In contrast, corticospinal neurons directly innervate motor neurons in primates to control fine movements. Spinal motor neurons also receive extensive projections from monoaminergic neurons from the brainstem, mostly from noradrenergic and serotonergic neurons. Many peptides are co-localized with monoaminergic projections, suggesting that they might be co-released with either serotonin or norepinephrine.

### 2.2 Abnormalities of Inputs to Motor Neuron in ALS

A number of cell types innervating motor neurons have been shown to degenerate in ALS animal models and/or patients. In this section, we will present the abnormalities by neurotransmitter.

#### 2.2.1 GABAergic Abnormalities

A number of GABAergic abnormalities have been documented in ALS.

Neuroimaging studies suggest that a loss of central nervous system inhibitory influence, in particular GABA, plays a role in ALS pathogenesis. Flumazenil binds to the benzodiazepine subunit of the GABA-A receptor present on neurons of the cerebral cortex. Using positron emission tomography (PET), a study showed a widespread loss of binding of [<sup>11</sup>C]-flumazenil in sporadic ALS patients. The abnormalities were less extensive in patients with SOD1-linked familial ALS patients, suggesting that GABAergic neurotransmission may be less severely impaired in these cases (Lloyd et al. 2000). Recent advances in proton magnetic resonance spectroscopy (<sup>1</sup>H-MRS) allow direct quantification of GABA in the cortex (Zhu et al. 2011). Using high resolution 3 Tesla edited <sup>1</sup>H-MRS, a study in a small number of subjects showed that decreased levels of GABA were present in the motor cortex of ALS patients compared to healthy controls (Foerster et al. 2012).

What could be the consequences of loss of cortical GABA? Studies with transcranial motor magnetic stimulation (TMS) suggest that the cortical hyperexcitability observed in ALS patients could be due to alterations in GABA-A signalling. In this technique, the intensity required to stimulate the motor cortex as well as the duration of the cortex silent period and the response to paired stimulation are measured, providing a good estimate of cortical excitability (Eisen and Weber 2001). The increased excitation observed in ALS patients may reflect a combination of an intrinsic hyperexcitability of the cortical inhibitory neurons. Although there are some conflicting data (Attarian et al. 2009), a loss of inhibition is suggested by several studies using paired-pulse stimulation that showed a decreased short-interval intracortical inhibition (SICI)

attributed to GABA-A receptors (Zanette et al. 2002; Yokota et al. 1996; Vucic and Kiernan 2006; Desiato et al. 1999; Caramia et al. 2000).

The role of cortical hyperexcitability in ALS remains debated, in particular with respect to its timing. Some authors have suggested it is an early feature (Mills and Nithi 1997; Prout and Eisen 1994; Caramia et al. 2000) whether others argued that it was a late phenomenon (Zanette et al. 2002). At least in SOD1-linked familial ALS, results obtained using a threshold-tracking TMS technique suggested that reduction of SICI is an early phenomenon that may precede the development of clinical symptoms (Vucic et al. 2008). Indeed, reduction of SICI was observed in three asymptomatic SOD1-mutation carriers who subsequently developed typical clinical features of ALS. A question is whether changes in excitability have implications on clinical symptoms. A study using threshold-tracking TMS showed a correlation between the reduction of SICI and fatigue (Vucic et al. 2011). The authors concluded that maladaptation of cortical processes related to degeneration of inhibitory GABAergic intracortical circuits may have a role in the development of fatigue in ALS.

All the previous results were obtained in cortical regions. It would be of high interest to obtain PET imaging or MRS data on inhibitory activity at the level of the spinal cord in ALS patients. However, technical limitations render such data currently inaccessible.

The GABAergic system has been poorly studied in animal models of ALS. In the few studies existing, GABAergic projections to motor neurons appeared much less affected than glycine interneurons (Chang and Martin 2009; Petri et al. 2005). However, the response of motor neurons to GABA might be slightly abnormal (Carunchio et al. 2008).

#### 2.2.2 Glycinergic Abnormalities

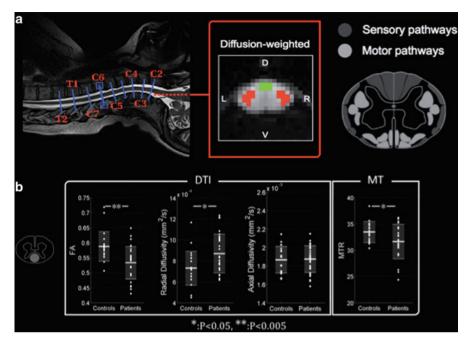
In patients, an impairment of Renshaw cells has been suggested by electrophysiological studies testing the mixed nerve silent period (Shefner and Logigian 1998) or using a conditioned H-reflex technique (Raynor and Shefner 1994). In contrast to spastic spinal cord-injured patients, spastic ALS patients showed strikingly reduced recurrent inhibition, suggesting that reduction of the recurrent inhibition suggested a different physiology for spasticity in ALS compared to spinal cord transection (Raynor and Shefner 1994). The role of Renshaw cells is however controversial in ALS since more distal muscles, more affected in ALS, lack recurrent inhibition (Mazzocchio and Rossi 2010).

The glycinergic system, and in particular the Renshaw cells, have been studied in more detail in experimental models of ALS. First, Renshaw cells lose connection with motor neurons in end-stage mice (Chang and Martin 2009; Wootz et al. 2012), and this might contribute to motor neuron hyperexcitability. Consistent with this, several authors described degeneration of glycinergic interneurons on spinal motor neurons of SOD1(G93A) mice and showed that glycinergic currents are decreased in SOD1(G93A) motor neurons (Chang and Martin 2009, 2011; Martin and Chang 2012). The pathology in glycinergic neurons appears however to follow the pathology of motor neurons, suggesting that it is either a very late event or a consequence of degeneration of motor neurons (Hossaini et al. 2011). A recent study in worms however suggested that, at least in this model, glycinergic interneurons are early and strongly affected by the disease process (McGown et al. 2012). More research is needed to ascertain this point in vertebrates.

#### 2.2.3 Glutamatergic Abnormalities

Glutamatergic abnormalities in ALS have been the subject of intense research in the last 30 years. Most research focused on abnormalities of glutamate release or reuptake by astrocytes as the source of increased glutamate in the cerebrospinal fluid (CSF). The contribution of neurons themselves has been scarcely studied. Direct glutamatergic innervation has been reported to be decreased in one study (Schutz 2005), but an opposite result has been recently published (Sunico et al. 2011). Consistent with an importance of glutamatergic innervation in motor neuron survival, decreasing vesicular glutamate transporter VGLUT2, responsible for a major part of glutamate release on motor neurons, decreased motor neuron degeneration, but did not have influence animals survival (Wootz et al. 2010). Although only indirectly innervating LMN in mice, it is clear that corticospinal motor neurons, a major glutamatergic population, degenerate in ALS mouse models (Ozdinler et al. 2011). It has been suggested a possible alteration of glutamatergic proprioceptive Ia afferents originating from muscle spindles primary endings, which have monosynaptic projections onto motor neurons. An impairment of the sensory pathways, with a degeneration of the dorsal funiculus, has been detected in animal model of SOD1-linked ALS (Guo et al. 2009). In a rat model of ALS, an impairment of dorsal roots was detected at an early and asymptomatic stage when motor neuron cell bodies did not appear to be compromised (Malaspina et al. 2010). Interestingly, sensory neurons that are functionally linked to muscle-spindle afferents by large-caliber myelinated fibers, corresponding to Ia fibers, were affected. The authors stated that these findings supported the pathogenic relevance in the development of the disease of sensory inputs coming from the muscles. Interestingly, two recent studies in a mouse model of another motoneuron degenerative disease, spinal muscular atrophy, demonstrated an early impairment of proprioceptive inputs to motor neurons. It was speculated that even a modest loss of proprioceptive function may have implications on motoneuron activity, since it has been estimated that one single Ia afferent can sprout collaterals and can develop over 2,700 synapses with up to 300 motoneurons in a motor neuron pool that innervates the medial gastrocnemius (Mendell and Henneman 1971).

Recently a spinal magnetic resonance imaging study showed that an impairment in the dorsal column, containing large myelinated sensory fibers, occurred at an early stage of the disease (Cohen-Adad et al. 2012) (Fig. 3). Together with findings in animal models, these in vivo findings in ALS patients suggest that the modifications glutamatergic input from type Ia muscle spindles to motoneurons may play a role in the pathogenesis of ALS.



**Fig. 3** Magnetic resonance imaging abnormalities of sensory pathways in ALS patients. (a) Regions of interest were selected in the lateral segment (turquoise blue) to include most of the lateral cortico-spinal tract and in the dorsal segment (purple) to include most of the dorsal columns (b) Individual plots of MRI metrics averaged in the dorsal columns. Differences were also detected in the lateral cortico-spinal tract (not shown). MTR stands for Magnetization Transfer Ratio.

#### 2.2.4 Serotonergic Abnormalities

Besides glutamatergic neurons, other long projection neurons targeting the motor neurons are affected by the disease, although this area is still largely unexplored. In particular, it was recently shown that serotonergic neurons profoundly degenerate in both patients and animal models. This degeneration is variable between nuclei and is mostly neuritic, with some loss of cell bodies. Most importantly, this degeneration contributes to spasticity in animal models (Dentel et al. 2012). Other consequences of serotonergic loss in ALS are still unexplored.

## 2.3 Strategies Targeting Glutamate

Multiple therapeutic strategies have been designed to target glutamate in ALS (Van Den Bosch et al. 2006). Indeed, glutamate levels are increased in the CSF of ALS patients, and motor neurons are exquisitely sensitive to increased glutamate because they express calcium permeable AMPA receptors, due to the very low levels of the GluR2 subunit on motor neurons and to abnormal editing of its mRNA (Kawahara and Kwak 2005; Shaw and Eggett 2000). This high permeability to

calcium, combined with poor buffering capacity of calcium in the cytoplasm of motor neurons, likely causes high cytosolic calcium increases in motor neurons in response to excess of glutamate, leading to cell death through excitotoxicity (Van Den Bosch et al. 2006). Multiple antiglutamatergic strategies have been built on this framework.

#### 2.3.1 Riluzole

Bensimon and collaborators demonstrated in 1994 that riluzole improved survival in ALS patients (Bensimon et al. 1994). A second larger confirmatory study published 2 years after this initial breakthrough confirmed this protective effect (Lacomblez et al. 1996a, b). Taking into account safety results, it was concluded from these two trials that the 100 mg dose of riluzole has the best benefit-to-risk ratio and that riluzole increases survival of patients with ALS. A meta-analysis of published riluzole trials in ALS has been recently performed and confirmed the increase in lifespan of ALS patients under riluzole (Miller et al. 2012). Taken together, interventional and population-based studies converge to demonstrate a consistent positive effect of riluzole on the survival of ALS patients. Apparently, this effect is stronger in patients treated early after symptom onset and in patients with bulbar onset. Although 3 months survival benefit is significant, there is no apparent slowing of disease progression. To our knowledge, riluzole is the only drug in the field of neurodegenerative diseases that proved increasing lifespan in multiple interventional trials. However, the underlying mechanism, long postulated to be glutamatergic, is extremely unclear, and multiple studies point to riluzole being rather an inhibitor of sodium persistent inward currents, rather than an antiglutamatergic drug (Bellingham 2011).

#### 2.3.2 Other Antiglutamatergic Agents

Results of therapeutic trials testing other antiglutamatergic agents have been disappointing. Gabapentin, which presumably acts through inhibition of glutamate biosynthesis, showed promising preclinical effects in SOD1 mice (Gurney et al. 1996), but a randomized clinical trial was negative in ALS patients (Miller et al. 2001). Lamotrigine, which is thought to inhibit glutamate release, also failed to demonstrate a beneficial effect in ALS patients (Ryberg et al. 2003). A potential therapeutic strategy to target excitotoxicity consists in the blockade of selected glutamate receptors. However, phase III clinical trials were negative in ALS when testing AMPA receptors blockers such as talampanel or topiramate (Cudkowicz et al. 2003). Clinical trial with NMDA receptor antagonists, dextromethorphan (Blin et al. 1996) or memantine (de Carvalho et al. 2010), was unsuccessful as well. The excitatory amino acid transporter EAAT2 is one of the major glutamate transporters expressed predominantly in astroglial cells and is responsible for 90 % of total glutamate uptake. The beta-lactam antibiotic ceftriaxone was shown to increase EAAT2 levels and to prevent motor neuron death in an animal model (Rothstein et al. 2005). However, a therapeutic trial recently failed to show a therapeutic effect in ALS patients (Cudkowicz 2012).

#### 2.4 Other Potential Strategies (Serotonin, Glycine)

To our knowledge, no drug based either on glycinergic transmission has been rigorously tested in an ALS clinical trial. Preclinical research is to be done. Concerning serotonergic transmission, serotonin levels are decreased in ALS blood (Dupuis et al. 2010), and decreased serotonin is responsible for the development of spasticity in ALS models (Dentel et al. 2012). Interestingly, providing ALS mice with serotonin precursor 5-hydroxytryptophan (5-HTP) increased their lifespan (Turner et al. 2003). However, here again, no drug targeting specific serotonin receptor subtype has been tested in models or patients.

# 3 Modulation of Motor Neuron Loss Through Increased Neurotrophic Activity

#### 3.1 Neurotrophic Factors: A Potential Therapeutic Strategy

It is widely recognized that providing more trophic support to dying motor neurons might slow down their degeneration. This is supported by a number of observations suggesting that neurotrophic factor expression and/or activity might be abnormal in ALS. For instance, expression of glial cell line-derived neurotrophic factor (GDNF), a major neurotrophic factor, is increased in ALS muscle (Lie and Weis 1998), as a likely consequence of denervation. Indeed, there are reported abnormalities in c-Ret, the GDNF receptor, in ALS motor neurons (Mitsuma et al. 1999; Ryu et al. 2011). Last, overexpressing GDNF is protective in multiple ALS models, using various mode of administration (Li et al. 2003, 2007 Manabe et al. 2002, 2003; Suzuki et al. 2008).

A second example is vascular endothelial growth factor (VEGF): VEGF expression is decreased in ALS spinal cord (Anagnostou et al. 2010) and in mouse models, and such a decrease is on its own sufficient to trigger ALS-like disease in mouse. Moreover, overexpression of VEGF is able to provide strong protection to animal models (Azzouz et al. 2004; Devos et al. 2004; Lambrechts et al. 2003; Oosthuyse et al. 2001; Storkebaum et al. 2005; Van Den Bosch et al. 2004; Wang et al. 2007).

Besides documented alterations in ALS, trophic factors are in general of interest in the treatment of ALS. Indeed, during development, motor neurons are produced in excess, and those that are unable to contact efficiently skeletal muscles are eliminated (Kanning et al. 2010). It is thus possible that similar mechanisms of survival might slow down degeneration of LMNs in adults affected with ALS. During development, survival of motor neurons is strictly under the control of the innervated limb through the production of multiple trophic factors that include GDNF or ciliary-derived neurotrophic factor (CNTF). During the last 20 years, a plethora of motor neurotrophic factors have been identified which are able in vitro to provide protection to isolated cultured motor neurons. Each of these neurotrophins is thus a potential candidate for ALS therapy.

#### 3.2 Treatment with Neurotrophic Factors in ALS Patients

Despite promising preclinical studies, direct administration of neurotrophic factor in human ALS patients failed to show a therapeutic effect in ALS patients. The major agent studied were brain-derived neurotrophic factor (BDNF) (BDNF Study Group 1999), CNTF (Miller et al. 1996), and insulin-like growth factor-1 (IGF-1) (Mitchell et al. 2007; Sorenson et al. 2008; Borasio et al. 1998; Lai et al. 1997). Clinical trial with xaliproden, an orally administered non-peptide compound that promotes the secretion of several endogenous motor neuron growth factors, failed to demonstrate a statistically significant therapeutic effect (Meininger et al. 2004). A human study with intraventricular injection of VEGF is currently in a phase I/II clinical trial.

#### 4 Conclusion

The last 20 years of research in the field of neuroprotection in ALS have been very successful for identifying potential strategies to improve the survival of motor neurons in ALS mouse models. However, none of these potential treatments successfully translated to the clinics. This failure might be due to multiple causes, and we would like to highlight two of these potential causes.

First, research efforts were almost exclusively devoted to identify compounds and/or strategies to provide protection to motor neuron cell body. However, such a protection is now known to be largely insufficient (Dupuis and Loeffler 2009). For instance, we and others have shown that providing a complete protection of motor neuron cell bodies does not ameliorate profoundly the phenotype of ALS mice because the remaining LMNs are still denervated and do not reinnervate affected muscles (Dewil et al. 2007; Gould et al. 2006; Rouaux et al. 2007). Thus, it is critical to identify the mechanisms that cause synapse loss, rather than motor neuron loss.

Second, despite it is largely assumed that ALS patients die of motor neuron degeneration, multiple events, many of them not occurring in motor neurons, strongly contribute to disease pathogenesis. In particular, despite weight loss is a strong predictor of survival, its causes remain unknown (Dupuis et al. 2011). The recent examples of microglial (Boillee et al. 2006) or serotonergic (Dentel et al. 2012) contributions to clinical picture should illuminate our need to understand how non-motoneuronal cell types could be targeted by specific treatments to achieve neuroprotection.

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# Neurotoxicity and ALS: Insights into Pathogenesis

# Steve Vucic and Matthew C. Kiernan

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### Abstract

Amyotrophic lateral sclerosis (ALS) is a rapidly progressive neurodegenerative disorder of the motor neurons in the spinal cord, brainstem, and motor cortex. While the mechanisms underlying the development of ALS remain to be fully elucidated, evidence is now emerging to suggest that the pathophysiological mechanisms underlying ALS are multifactorial, reflecting a complex interaction between causal genes and local environment. In particular, dysfunction of metabolic pathways including excessive oxidative stress, glutamate excitotoxicity, mitochondrial dysfunction, and defective axonal transport

M.C. Kiernan (🖂)

S. Vucic

Sydney Medical School Westmead, University of Sydney, Sydney, Australia

Neuroscience Research Australia, Randwick, Sydney, NSW, Australia e-mail: s.vucic@neura.edu.au

Bushell Professor of Neurology, Brain and Mind Research Institute, University of Sydney, Sydney, Australia

Neuroscience Research Australia, Randwick, Sydney, NSW, Australia e-mail: m.kiernan@unsw.edu.au

systems along with abnormalities of nonneuronal supporting cells may cause critical injury to target proteins and organelles within the motor neuron, thereby leading to neurotoxicity and neurodegeneration in ALS. The clinical effectiveness afforded by anti-glutamatergic agents such as riluzole underscores the importance of glutamate excitotoxicity in the development of neurodegeneration in ALS, with anterior horn cell degeneration mediated by corticomotoneuronal hyperexcitability via an anterograde transsynaptic process. This chapter will review current understanding of the pathophysiological mechanisms underlying the development of ALS, with a particular focus on the role of neurotoxicity in mediating neurodegeneration in ALS.

### **Keywords**

ALS • Glutamate excitotoxicity • Neurodegneration • Neurotoxicity

### 1 Introduction

Amyotrophic lateral sclerosis (ALS) is a rapidly progressive neurodegenerative disorder of motor neurons in the spinal cord, brainstem, and motor cortex (Kiernan et al. 2011). The ALS phenotype is clinically characterized by the presence of upper and lower motor neuron abnormalities (Kiernan et al. 2011; Vucic et al. 2007; Winhammar et al. 2005), whereby lower motor neuron (LMN) features include fasciculations, muscle wasting, and weakness, while upper motor neuron features include weakness, slowness of movement, increased tone, and hyperreflexia. This unique combination of upper and lower motor neuron abnormalities, not encountered in other neurodegenerative diseases, led Charcot to postulate the mechanism of neuronal involvement in ALS pathogenesis (Charcot and Joffroy 1869). Despite Charcot's initial observations, the precise pathophysiological mechanisms, and even the site of disease onset, remain the subject of ongoing debate (Ravits et al. 2007).

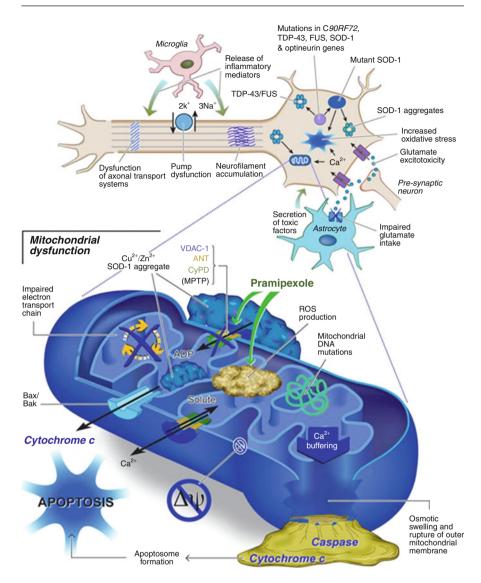
Evidence is now emerging to suggest that the pathogenic mechanisms underlying ALS are multifactorial, reflecting interaction between causal genes and local environment (Boillee et al. 2006a; Kiernan et al. 2011; Vucic and Kiernan 2009; Winhammar et al. 2005). These complex pathways, including oxidative stress, glutamate excitotoxicity, mitochondrial dysfunction, and defective axonal transport systems (Fig. 1), combined with abnormalities of nonneuronal supporting cells such as the astrocytes, may cause injury of critical target proteins and organelles within the motor neuron, thereby resulting in neurotoxicity and degeneration in ALS (Gonzalez de Aguilar et al. 2007; Gros-Louis et al. 2006; Haidet-Phillips et al. 2011; Neusch et al. 2007; Pasinelli and Brown 2006; Patel and Maragakis 2002; Vucic and Kiernan 2009). As such, this chapter will review current understanding of the pathophysiological mechanisms underlying the development of ALS, with a particular focus on the role of neurotoxicity.

## 1.1 Glutamate-Mediated Excitotoxicity

Glutamate-mediated excitotoxicity appears to be an important mechanism in ALS pathogenesis (Boillee et al. 2006; Kiernan et al. 2011). Glutamate is the major excitatory neurotransmitter in the central nervous system (Heath and Shaw 2002; Watkins and Evans 1981), synthesized from reductive deamination of alpha-ketoglutarate or from the action of amino acids of aminotransferases (Heath and Shaw 2002). Approximately 20% of the total glutamate pool is stored in presynaptic nerve terminals, and during impulse transmission, glutamate is released from presynaptic neurons through the effects of depolarization, diffusing across the synaptic cleft to activate postsynaptic receptors. Their excitatory signal is terminated upon removal of glutamate from the synaptic cleft by specific glutamate reuptake transporters located on both neurons and astrocytes (Dong et al. 1999; Vandenberg 1998). Within presynaptic astrocytes, glutamate is converted into glutamine by the enzyme glutamine synthetase and then returned to the neuron for resynthesis of glutamate (Laake et al. 1995).

Glutamate receptors are broadly classified into ionotropic or metabotropic receptors (Heath and Shaw 2002). Binding of glutamate to its ionotropic receptors results in a conformational change within the receptor, thereby enabling a passage of Na<sup>+</sup> and Ca<sup>2+</sup> ions through a central pore. Metabotropic glutamate receptors are linked via G-proteins to second-messenger enzymes, which in turn can regulate a host of cellular activities (Simeone et al. 2004). Based on pharmacological studies, glutamate ionotropic receptors are further classified as (i) N-methyl-Daspartate (NMDA), (ii)  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazoleproprionic acid (AMPA), and (iii) kainite receptors. This pharmacological classification is supported by cloning studies that have identified six different families of glutamate ionotropic receptors that conform to the original agonist studies (Simeone et al. 2004). N-methyl-D-aspartate receptors are permeable to influx of  $Na^+$  and  $Ca^{2+}$  and efflux of K<sup>+</sup> (Simeone et al. 2004). An essential feature of NMDA receptors is their voltage-dependent blockade by Mg<sup>2+</sup> binding within the channel pore, which can be alleviated by depolarization (MacDermott et al. 1986). NMDA receptors are involved in excitatory neurotransmission, which is characterized by a slow rise time and decay. As such, the NMDA receptors are involved in complex physiological processes, such as generation of rhythmic motor activity (Traven et al. 1993), regulation of neuronal migration during embryogenesis (Komuro and Rakic 1993), and memory (Bliss and Collingridge 1993).

The NMDA receptor complex is composed of different subunits derived from 6 genes: NMDAR1 (eight splice variants described), NMDAR2 (A–D), and NMDAR3 (A,B) (Heath and Shaw 2002; Simeone et al. 2004). While the NMDAR1 subunit forms the basic structure of the receptor (Heath and Shaw 2002), the NMDAR2 subunit determines ion channel properties and forms ligandbinding sites (Kutsuwada et al. 1992; Meguro et al. 1992; Michaelis 1998). Functional and pharmacological properties of NMDA receptors are determined through specific combination of NMDAR1 and NMDAR2 subunits (Kutsuwada et al. 1992;



**Fig. 1** Multiple interacting pathophysiological mechanisms appear to underlie the development of amyotrophic lateral sclerosis (*ALS*). Glutamate-mediated excitotoxicity is an important mechanism ALS progression and seems to precede the clinical development of ALS. Dysfunction of the excitatory amino acid transporter type 2 (EAAT2), located on astrocytes, in part induces the development of glutamate excitotoxicity via excessive extracellular accumulation of glutamate. In addition, activation of nonneuronal cells (astrocytes and microglia) in ALS results in secretion of proinflammatory cytokines and other cytotoxic factors that ultimately results in further neurotoxicity and degeneration. In conjunction with glutamate-mediated excitotoxicity, other molecular processes induce neurotoxicity via multifactorial mechanisms. Within the neuron, mutations in a host of ALS-related genes, including C9orf72, superoxide dismutase 1 (*SOD1*), TDP-43, and FUS, result in ALS via multiple mechanisms. Specifically, mutations in the SOD-1

Monyer et al. 1992). In addition, there are regional variations in the expression of NMDA receptor subtypes (Ciabarra and Sevarino 1997; Ishii et al. 1993; Kutsuwada et al. 1992; Monyer et al. 1992; Watanabe et al. 1993a, b, 1994a, b), with the NMDAR3B subunit heavily expressed in somatic motoneurons (Chatterton et al. 2002; Nishi et al. 2001).

AMPA receptors mediate a rapid influx of monovalent ions, such as Na<sup>+</sup>, K<sup>+</sup>, and chloride (Cl<sup>-</sup>) but unlike NMDA receptors are impermeable to Ca<sup>2+</sup> (Heath and Shaw 2002). Four AMPA receptor subtypes have been cloned (*Glu*R1-4) and are composed of three transmembrane domains (M1, M3, M4) and a fourth cytoplasmic hairpin loop (M2), which contributes to the pore-lining region (Dingledine et al. 1999; Simeone et al. 2004). The AMPA receptor exists as a pentameric structure in vivo, which is formed by the arrangement of subunits to create receptor diversity (Heath and Shaw 2002). The GluR2 subunit influences the Ca<sup>2+</sup> permeability of AMPA receptors, whereby those receptors expressing an immature GluR2 subunit are more permeable to Ca<sup>2+</sup> results in neurodegeneration through activation of Ca<sup>2+</sup>-dependent pathways (Heath and Shaw 2002; Simeone et al. 2004).

# 1.2 The Role of Glutamate in ALS Pathogenesis

As discussed, glutamate excitotoxicity is mediated by excessive activation of postsynaptic glutamate receptors (Heath and Shaw 2002). In ALS, glutamate excitotoxicity has been postulated to induce anterior horn cell degeneration via a transsynaptic anterograde process mediated by corticomotoneurons (Eisen et al. 1992). Support for such a mechanism has been provided by transcranial magnetic stimulation studies (TMS) which have demonstrated that cortical hyperexcitability, a biomarker of glutamate excitotoxicity, is an early feature in sporadic and familial ALS, linked to motor neuron degeneration (Blair et al. 2010; Caramia et al. 1991; Desiato et al. 2002; Eisen et al. 1993; Prout and Eisen 1994; Vucic and Kiernan 2006, 2009, 2010; Vucic et al. 2008). In addition, longitudinal studies in asymptomatic SOD-1 mutation carriers revealed that cortical hyperexcitability developed prior to the clinical onset of ALS (Vucic et al. 2008), a feature also evident in the G93A SOD-1 mouse model (Browne et al. 2006). Of relevance, loss of  $\gamma$ -aminobutyric acid (GABA) secreting parvalbumin-positive inhibitory interneurons in the motor cortex of ALS patients may further contribute

**Fig. 1** (continued) gene result in toxic gain of function of the SOD1 enzyme which affects a host of critical cellular organelles, such as DNA/RNA metabolism. In addition, mitochondrial dysfunction is a feature of ALS, linked to glutamate excitotoxicity and SOD-1 gene mutations, resulting in a reduced production of ATP and calcium-sequestering ability, as well as an increase in free radical formation. Of further relevance, mitochondrial dysfunction may contribute to glutamate excitotoxicity. Ultimately, these multiple pathogenic processes result in critical cell dysfunction and motor neuron degeneration

to the development of cortical hyperexcitability (Nihei et al. 1993), a finding underscored by recent neuroradiological studies reporting a significant reduction of GABA within the motor cortex of ALS patients (Foerster et al. 2012).

Molecular-based studies have provided further evidence for glutamate-mediated excitotoxicity in ALS. Specifically, molecular studies established significant reduction in the expression and function of the astrocytic glutamate transporter (EAAT2), which mediates glutamate reuptake at synapses thereby reducing glutamate excitotoxicity, in the superoxide dismutase-1 (SOD-1) mouse model and the motor cortex and spinal cord of ALS patients (Boillee et al. 2006; Ionov 2007; Rothstein et al. 1993, 1995, Trotti et al. 1999). Of further relevance, dysfunction of the EAAT2 transporter appears to be a preclinical feature in the SOD-1 mouse model (Boston-Howes et al. 2006; Gibb et al. 2007), and an increase in the expression and transporter activity of EAAT2 increases the life span of mutant SOD-1 mice (Rothstein et al. 2005).

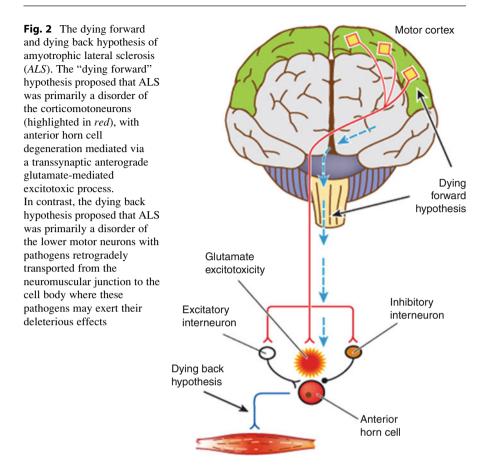
At a postsynaptic level, increased expression of AMPA receptors with the unedited GluR2 subunit has been reported in ALS (Kawahara et al. 2004; Kwak and Kawahara 2005; Takuma et al. 1999; Van Damme et al. 2002, 2005). This editing defect appears to be specific for ALS, thereby rendering the motor neurons more permeable to Ca<sup>2+</sup>, potentially explaining the increased sensitivity of motor neurons to excitotoxicity (Cox et al. 2007; Heath and Shaw 2002). Further support for glutamate excitotoxicity has been provided by the clinical benefit of riluzole in ALS patients (Bensimon et al. 1994; Cheah et al. 2010; Gurney et al. 1996, 1998; Lacomblez et al. 1996). Specifically, riluzole is a glutamate antagonist that exerts effects in the central nervous system by reducing the release of glutamate from presynaptic nerve terminals and enhancing the reuptake of glutamate (Azbill et al. 2000; Cheah et al. 2010; Quinlan 2011; Wang et al. 2004).

For the glutamate hypothesis to be a plausible mechanism of motor neuron degeneration in ALS, it must explain how motor neurons became selectively damaged through overactivity of the glutaminergic system and provide a mechanism by which degeneration occurs. A number of cell-specific molecular features possessed by the motor neurons render them vulnerable to glutamate toxicity in ALS. Motor neurons affected in ALS preferentially express AMPA receptors lacking the functional GluR2 subunit, thereby rendering the motor neurons more permeable to Ca<sup>2+</sup> (Kawahara et al. 2004; Kwak and Kawahara 2005; Van Damme et al. 2002, 2005). In addition, motor neurons vulnerable to degeneration lack the intracellular expression of proteins parvalbumin and calbindin D28k which are required to buffer intracellular Ca<sup>2+</sup> (Ince et al. 1993). Of further relevance, increased expression of the inositol 1.4,5triphosphate receptor 2 (ITPR2) gene was reported in ALS (van Es et al. 2007). The ITPR2 is involved in glutamate-mediated neurotransmission, whereby stimulation of glutamate receptors results in binding of inositol 1.4, 5-triphosphate to ITPR2, which subsequently increases intracellular calcium (Choe and Ehrlich 2006; van Es et al. 2007). Aberrant activity of ITPR2 results in higher intracellular concentration of Ca2+ leading ultimately to neurodegeneration (Gutstein and Marks 1997). Of further relevance, motor neurons in ALS, at least in animal models, appear to be larger, with an increase in distal dendritic branching (Amendola and Durand 2008). Consequently, the input conductance of the motor neurons is increased rendering the motor neuron more vulnerable to electrical and metabolic stresses, in particular those imparted by glutamate excitotoxicity (Quinlan 2011).

Although details of the molecular mechanisms by which glutamate exerts neurotoxicity are still to be fully elucidated, several pathways have been defined. Initially, an influx of Na<sup>+</sup> and Cl<sup>-</sup> ions occurs along with water molecules, resulting in acute neuronal swelling that is reversible with removal of agonist (Choi 1987; Shaw and Kuncl 2002; Stys 1998). Subsequently, an influx of Ca<sup>2+</sup> ions occurs via activation of ionotropic receptors such as the NMDA and Ca<sup>2+</sup>-permeable AMPA receptors, as well activation of voltage-gated Ca<sup>2+</sup> channels (Choi 1987; Miller et al. 1989). Ultimately, activation of these ionic pathways results in increased intracellular Ca<sup>2+</sup> concentration and activation of Ca<sup>2+</sup>-dependent enzymatic pathways leading to neuronal death (Cox et al. 2007; Meldrum and Garthwaite 1990; Regan et al. 1995; Shaw and Kuncl 2002). Further, glutamate excitotoxicity results in production of free radicals that can further damage the intracellular organelles to thereby cause cell death (Bondy and Lee 1993; Lees 1993; Maher and Davis 1996).

In his original manuscript, Charcot concluded that ALS was a disorder of the brain and that lower motor neuron degeneration resulted from downstream effects (Charcot and Joffroy 1869). This view was not universal, and some of Charcot's contemporaries, such as Gower's, argued that upper and lower motor neuron degeneration were independent. Over the past two decades the site of ALS onset has been revisited to a large extent precipitated by the advent of modern noninvasive technologies such as TMS. Three schools of thought have developed pertaining to the role of the UMN and related pathophysiological processes in ALS: (i) "the *dying forward*" hypothesis; (ii) "the *dying back*" hypothesis, and (iii) "the *independent degeneration*" hypothesis (Fig. 2).

The dying forward hypothesis proposed that ALS was primarily a disorder of the corticomotoneurons, which connect monosynaptically with anterior horn cells (Eisen et al. 1992). Corticomotoneuronal hyperexcitability was postulated to induce anterior horn cell degeneration transsynaptically via an anterograde glutamatemediated excitotoxic process (Eisen et al. 1992; Vucic and Kiernan 2009). This dying forward hypothesis was based on a number of poignant clinical observations including (i) relative preservation of extraocular and sphincter muscles in ALS, postulated to be due to a paucity of corticomotoneuronal projections onto the motor nuclei innervating these muscles; (ii) absence of an animal model of ALS, ascribed to a lack of direct corticomotoneuronal-anterior horn cell connections (Armand 1982; Lemon and Griffiths 2005); (iii) rarity of pure lower motor neuron forms of ALS, with subclinical upper motor neuron dysfunction invariably detected with TMS studies (Eisen and Shtybel 1990); and (iv) the specificity of dissociated muscle atrophy (Eisen and Kuwabara 2012; Kuwabara et al. 2008; Menon et al. 2011; Wilbourn 2000), termed the split hand phenomenon in ALS remains best explained by a dying forward mechanism (Eisen and Kuwabara 2012; Menon et al. 2011).



Utilizing TMS technology, it is becoming increasingly apparent that cortical hyperexcitability develops as an early feature in sporadic and familial ALS, linked to the process of motor neuron degeneration (Caramia et al. 1991; Desiato et al. 2002; Eisen et al. 1993; Prout and Eisen 1994; Vucic and Kiernan 2006, 2009, 2010; Vucic et al. 2008). Furthermore, longitudinal studies in asymptomatic SOD-1 mutation carriers established that cortical hyperexcitability developed prior to the clinical onset of ALS (Vucic et al. 2008), a feature also reported in the G93A SOD-1 mouse model (Browne et al. 2006). In keeping with a cortical origin of ALS is the now accepted view that ALS and frontotemporal dementia (FTD) represent an overlapping continuum of the same disorder (Lillo and Hodges 2009; Neumann et al. 2006), an observation further underscored by recent genetic discoveries that increased hexanucleotide repeat expansions in the first intron of C9ORF72 gene (9p21) was associated with both ALS and FTD (DeJesus-Hernandez et al. 2011; Renton et al. 2011; Traynor 2012).

The dying back hypothesis proposed that ALS is primarily a disorder of the lower motor neurons, with pathogens retrogradely transported from the

neuromuscular junction to the cell body where they exert their deleterious effects (Chou and Norris 1993). Although some pathological studies have indirectly supported a dying back process (Gould et al. 2006; Pagani et al. 2006; Pun et al. 2006), no pathogens of any type has been identified in relation to ALS. The presence of widespread dysfunction within the frontal cortex, including the primary, supplementary, and prefrontal motor cortices in ALS, remains difficult to reconcile with any dying back process (Miller et al. 2009; Turner et al. 2009; Vucic et al. 2007). In addition, the absence of central pathology in other lower motor neuron disorders such as Kennedy's disease or poliomyelitis provides a further argument against a dying back process (Eisen and Weber 2001; Vucic and Kiernan 2008).

The independent degeneration hypothesis suggested that the upper and lower motor neurons degenerated independently (Gowers 1886–1888). Limited neuropathological studies provided indirect support for independent degeneration, whereby the degeneration of upper and lower motor neurons appeared to be independent (Kiernan and Hudson 1991; Pamphlett et al. 1995). These correlative morphological techniques, however, were significantly confounded by the anatomical and functional complexity of the corticomotoneuronal system (Flament et al. 1993). In particular, there remains considerable variability in the corticomotoneuronal to anterior horn cell ratio, due to synaptic changes, and as such, attempts to correlate upper and lower motor neurons as a "one-off" on autopsy studies are divorced from clinical and in vivo reality (Eisen and Weber 2001).

## 1.3 Neurotoxicity and the Role of SOD-1 Gene

Mutations in the SOD-1 gene, the first ALS gene reported and mapped to the long arm of chromosome 21 [21 q22.1] (Siddique et al. 1991), was postulated to exert pathogenic effects by resulting in acquisition of aberrant cytotoxic enzyme activity (Andersen 2006a; Dewil et al. 2004; Robberecht 2002). The SOD-1 gene spans 11 kilobases of genomic DNA, comprises five exons and four introns (Levanon et al. 1985), and encodes a highly conserved 153-amino acid long protein, which together with a catalytic copper (Cu) ion and a stabilizing zinc (Zn) ion form a subunit (Andersen 2006; Levanon et al. 1985). A disulfide bridge stabilizes each subunit, and the two identical subunits combine through non-covalent bonds to form the Cu-Zn SOD-1 enzyme. The main function of the SOD-1 enzyme involves free radical scavenging whereby the enzyme catalyses the conversion of the superoxide anion to molecular oxygen and hydrogen peroxide, which in turn is reduced to water by glutathione peroxidase and catalase (Andersen 2006; Fridovich 1986; Shaw and Kuncl 2002). The Cu-Zn SOD enzyme constitutes 0.5-1% of soluble protein in the brain and spinal cord and is located within the cytosol and nucleus and between two mitochondrial membranes (Andersen 2006; Bowling et al. 1995; Pardo et al. 1995).

To date, over 150 different mutations have been reported in the SOD-1 gene, with the majority of being missense mutations, resulting in changes in single amino acids but preserving the SOD-1 protein length. The remaining mutations are either nonsense or deletion mutations that either introduce novel nucleotides or remove existing nucleotides resulting in alteration of the polypeptide length (Andersen 2006; Dewil et al. 2004; Gros-Louis et al. 2006). An autosomal dominant pattern of inheritance is evident with most mutations, except for the D90A mutation, which may be transmitted in an autosomal recessive manner. Compound heterozygosity has also been reported with SOD1 FALS, whereby two siblings with a slowly progressive ALS phenotype may be carriers of both the D90A and D96N mutations (Hand et al. 2001). The mutations are widely distributed throughout the gene with preponderance for exon 4 and 5 (Andersen 2006; Andersen et al. 2003; Cudkowicz et al. 1997; Radunovic and Leigh 1996). Globally, the most frequent mutation is the substitution of aspartate for alanine (D90A), followed by alanine to valine (A4V) and isoleucine for threonine (I113T) (Andersen 2006; Dewil et al. 2004).

Evidence for a neurotoxic gain of function of the SOD-1 enzyme has been provided by several lines of evidence. Specifically, increased activity of the SOD-1 enzyme was reported in the transgenic SOD-1 mouse models, supporting a toxic gain of function mechanism (Bruijn et al. 1997; Gurney et al. 1994; Ripps et al. 1995; Wong et al. 1995). Of further relevance, SOD-1 knockout mice failed to develop the ALS phenotype, thereby suggesting the importance of SOD-1 enzyme activity in ALS pathogenesis (Reaume et al. 1996). Aberrant biochemical activity of the SOD-1 enzyme has been reported to underlie this toxic gain of function (Andrus et al. 1998; Beckman et al. 1993; Bruijn et al. 1997, 1998, 2004). Specifically, SOD-1 gene mutations reportedly induced structural changes in the SOD-1 enzyme, enabling substrates other than the superoxide anion to gain access to the active center, thereby resulting in increased production of hydroxyl and free radicals (Bogdanov et al. 1998; Liu et al. 1998). In addition, the mutated SOD-1 enzyme also accepts peroxynitrate as a substrate, resulting in nitration of tyrosine residues on critical cellular proteins (Beckman et al. 1993; Beckman and Koppenol 1996; Crow et al. 1997). Ultimately, this aberrant biochemical activity of the SOD-1 enzyme resulted in cell injury and death (Beckman et al. 1993). Importantly, upregulation of protein-tyrosine nitration has been reported in anterior horn cells (Abe et al. 1997; Chou et al. 1996) as well as elevation of 3-nitortyrosine in the spinal cord of ALS patients and transgenic SOD-1 mice (Beal et al. 1997; Ferrante et al. 1997).

Mutations in the SOD-1 gene may also result in improper binding of zinc to the mutated SOD-1 peptide, thereby allowing reduction of SOD-1 bound copper, which in turn results in formation of superoxide anions and cell injury (Estevez et al. 1999). Diminished metal ion binding by the mutated SOD-1 peptide may also release zinc and copper ions, thereby further contributing to neurotoxicity and degeneration (Pasinelli and Brown 2006). Of further relevance, aberrant SOD-1 enzyme activity may result in oxidative stress via upregulation of proinflammatory cytokines (Hensley et al. 2006). Specifically, increased expression of proinflammatory mediators such as nitric oxide, interleukins 1, 6, and 12 has been reported in the SOD-1 mouse model with resultant neurotoxicity to motor neurons in spinal cord preparations (Kim et al. 2006). Importantly, antagonism of these agents with neutralizing antibodies resulted in increased motor neuron survival (Kim et al. 2006).

Alternatively, mutations in the SOD-1 gene may lead to conformational instability and misfolding of the SOD-1 peptide, resulting in formation of toxic

intracellular aggregates. In the transgenic SOD-1 mouse model and human ALS cases, immunoreactive SOD-1 aggregates were reported in motor neurons and glial cells (Bruijn et al. 1998; Jonsson et al. 2006; Zetterstrom et al. 2007). Whether the intracellular aggregates were neurotoxic to motor neurons remains unknown, although a number of possible cytotoxic mechanisms have been proposed, including (i) co-aggregating with vital cellular constitutes, (ii) inhibiting normal proteosome function, and (iii) exerting mechanical or biochemical effects on the cell, such as disruption of axonal transport systems (Bruijn et al. 1997, 1998; Pasinelli and Brown 2006; Williamson and Cleveland 1999).

### 1.4 Mitochondrial Dysfunction

In conjunction with glutamate excitotoxicity and oxidative stress, there is mounting evidence that mitochondrial dysfunction exerts an important role in the pathophysiology of ALS (Boillee et al. 2006; Chung and Suh 2002; Higgins et al. 2003; Kirkinezos et al. 2005; Lederer et al. 2007; Pasinelli and Brown 2006; Xu et al. 2004). Mitochondria are intracellular organelles whose main function is to generate energy for the cell in the form of ATP. Under conditions of excessive  $Ca^{2+}$  load, as may be evident with glutamate excitotoxicity (Dugan and Choi 1994), mitochondrial production of free radicals induces injury of critical neuronal cellular proteins and DNA. In addition, mitochondria remain sensitive to free radical damage at both the protein and DNA level, resulting in further mitochondrial dysfunction (Bowling and Beal 1995). Mitochondrial damage may in turn enhance glutamate excitotoxicity by disrupting the normal resting membrane potential, thereby resulting in a loss of the normal voltage-dependent Mg<sup>2+</sup>-mediated block of NMDA receptor channels (Heath and Shaw 2002; Shaw and Kuncl 2002).

Mitochondrial degeneration and dysfunction has been reported in ALS patients and in the transgenic SOD-1 mouse model (Higgins et al. 2003; Kong and Xu 1998; Xu et al. 2004). Ultrastructural abnormalities of muscle mitochondria, paracrystalline inclusions, and abnormal cristae have been reported in ALS (Chung and Suh 2002; Comi et al. 1998; Lederer et al. 2007). Dysfunction of mitochondrial enzymes involved in energy generation, such as cytochrome C oxidase and respiratory chain complexes I and IV, as well as downregulation of nuclear genes encoding mitochondrial components within the motor cortex has been reported in ALS (Comi et al. 1998; Fujita et al. 1996; Jung et al. 2002; Kirkinezos et al. 2005; Lederer et al. 2007). Of further relevance, mitochondrial dysfunction including reduction in protein import, impairment in Ca<sup>2+</sup> sequestering ability, and an exaggerated depolarizing response of the inner mitochondrial membrane to Ca<sup>2+</sup> stimulation may occur in the presymptomatic stages of ALS (Bilsland et al. 2008; Damiano et al. 2006; Jaiswal et al. 2009; Li et al. 2010; Nguyen et al. 2009). Ultimately, severe damage to the mitochondrial membrane potential, respiration, and electron transfer chain ensues, resulting in reduced ATP synthesis and neurodegeneration (Quinlan 2011).

The transportation and distribution of mitochondria within the neurons appears to be impaired in ALS (Ouinlan 2011). Mitochondria are normally highly mobile organelles, evident in both the axons and dendrites (MacAskill et al. 2010). The movement of mitochondria is regulated through Ca<sup>2+</sup> signalling and synaptic activity (MacAskill et al. 2010). An increase in intracellular Ca<sup>2+</sup> concentration, as occurs with glutamate excitotoxicity, interrupts the movement of mitochondria within the cell, in particular at the level of the synapse (MacAskill et al. 2009). Abnormalities of mitochondrial distribution and transport have been reported in ALS, with evidence of reduced distribution in the axons and more frequent pauses in mitochondrial movements (Bilsland et al. 2010). Importantly, the slow and fast axonal transport systems, vital for mitochondrial transport, seem to be impaired in ALS and have been linked to glutamate excitotoxicity (Bilsland et al. 2010; De Vos et al. 2007; Quinlan 2011). Ultimately, this interruption in mitochondrial mobility may result in depletion of energy supply in critical neuronal segments, essential for the maintenance of the resting membrane potential and generation of action potentials, with resultant neuronal degeneration.

From a therapeutic perspective, a recent phase II trial of dexpramipexole, a pharmacological agent that enhances mitochondrial function (Cheah and Kiernan 2010), was shown to be effective in slowing ALS disease progression and reducing mortality over a 24-week period (Cudkowicz et al. 2011). Currently, a phase III, multicenter international trial is underway to assess the clinical efficacy of dexpramipexole as add-on therapy to riluzole in ALS (ClinicalTrials.gov-NCT01281189).

# 2 Non-neuronal Cells and Neurotoxicity in ALS

Dysfunction of nonneuronal cells, astrocytes, and microglia also appears to be important in ALS pathogenesis (Boillee et al. 2006a, b; Haidet-Phillips et al. 2011; Neusch et al. 2007). Molecular studies in the SOD-1 mouse model have established that expression of the mutant SOD-1 protein in either the motor neurons or astrocytes in isolation failed to induce degeneration, thereby suggesting that the pathogenic process in ALS may involve a complex interaction between motor neurons interacting with nonneuronal cells (Gong et al. 2000; Lino et al. 2002; Pramatarova et al. 2001). Underscoring the importance of nonneuronal cells in ALS pathogenesis are findings that motor neuron toxicity appears to be modulated by expression of mutant SOD-1 in nonneuronal cells (Clement et al. 2003). However, mutant SOD-1 expressing nonneuronal cells seem to be more involved in the regulation of disease progression rather than onset of ALS (Beers et al. 2006; Boillee et al. 2006).

Similar to animal models, nonneuronal cells seem also to be an important pathogenic factor in the human disease. Specifically, astrocytes derived from postmortem spinal cord neural progenitor cells (NPCs) of sporadic and familial (SOD1) patients were selectively toxic to motor neurons (Haidet-Phillips et al. 2011). Upregulation of inflammatory genes encompassing chemokines, proinflammatory cytokines and components of the complement cascade, was evident in the neurotoxic astrocytes, further suggesting that neurotoxicity may in

part be mediated by an inflammatory mechanism. In addition, downregulation of mutant and wild-type SOD-1 expression in astrocytes derived from familial and sporadic ALS, respectively, was neuroprotective (Haidet-Phillips et al. 2011). Taken together, these findings suggest that astrocyte-mediated neurotoxicity contributed to neurodegeneration in familial (SOD-1) and sporadic ALS, thereby suggesting that novel cell-based approaches, particularly focusing on nonneuronal supportive cells, may be therapeutic in ALS.

# 3 Conclusion

Multiple interacting molecular and genetic mechanisms appear to underlie the development of motor neurodegeneration in ALS. Studies in animal models and human ALS have suggested an important role for glutamate neurotoxicity in ALS pathogenesis. Specifically, neurophysiological studies have suggested that cortical hyperexcitability, a biomarker of glutamate neurotoxicity, appears as a primary event in ALS, with motor neuron degeneration mediated via transsynaptic anterograde mechanisms. In conjunction with glutamate excitotoxicity, other molecular processes including mitochondrial dysfunction and abnormalities of axonal transport, together with neurotoxicity of the mutant SOD-1 enzyme, oxidative stress, and dysfunction of nonneuronal supporting cells, such as astrocytes, appear to contribute to ALS pathogenesis. From a therapeutic perspective, further insights into ALS pathogenesis will undoubtedly result in development of novel therapeutic strategies.

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# Neurotoxicity and Neuroprotection in Spinal Cord Injury

# Renée Morris

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### Abstract

In the present chapter, the members of the neurotrophin family of neurotrophic factors and their receptors are first presented. This first section is then followed by a description of the neurotrophin-mediated signalling pathways that regulate cell survival, with particular emphasis on the "signalling endosome model" of intracellular signalling. The main functions of these neurotrophin-mediated signalling pathways are then described, and evidence that neurotrophic factors have the capacity to protect spinally projecting neurons against cell death and atrophy is subsequently examined. In this context, the neurotrophin-associated molecular

### R. Morris

Translational Neuroscience Facility, School of Medical Sciences, The University of New South Wales, Sydney, NSW, Australia

e-mail: renee.morris@unsw.edu.au

cascades of events that lead to the regulation of cell survival are presented. The protective role of the different neurotrophic factors for injured spinal cord-projecting neurons and their capacity to upregulate regeneration-associated genes is subsequently examined. The regenerative property of neurotrophins on lesioned sensory and motor axons is then introduced, and evidence that the delivery of neurotrophic factors, if available in sufficient concentration, can outweigh the effect of inhibitory molecules present in the scar tissue is reviewed. Finally, the different strategies currently under investigation to deliver neurotrophin to the injured spinal cord are reviewed. Here, special attention is given to the evidence for and against a potential role for neurotrophins in the recovery of motor function.

#### Keywords

Motor function • Neurotrophic factors • Neurotrophin • Regeneration • Spinal cord injury • Spinal cord-projecting neurons

# 1 Introduction

The propensity of the adult spinal cord to regenerate is only limited, and damage that disrupts the long myelinated fiber pathways within the cord results in permanent functional deficits. Over the last three decades, research in neuroscience has identified a set of pathophysiological events that occur within the injured spinal cord tissue and that contribute to halt axonal regeneration. One of the obstacles to axonal growth through the site of injury is the presence, in the glial scar, of potent growth inhibitory extracellular matrix molecules such as chondroitin and keratan sulfate proteoglycans (for a review, see Bradbury and Carter 2011). Another major impediment to axonal regeneration through traumatic damage to the cord is the existence of the myelin-associated axon growth inhibitory molecules Nogo-A, myelin-associated glycoprotein (MAG), and oligodendrocyte myelin glycoprotein (OMgp) (recently reviewed by Sharma et al. 2012). However, the presence of these inhibitory molecules within the damaged tissue is only part of the reason why injured axons fail to regenerate through a spinal cord injury. Indeed, a large population of spinal cord-projecting neurons that are axotomized after a spinal cord injury have been shown to exhibit a downregulation of genes that are directly involved in axonal regeneration (e.g., GAP-43 and T $\alpha$ 1 Tubulin), and this change in gene expression coincides with neuronal death and atrophy (Tetzlaff et al. 1991).

In the developing central nervous system (CNS), where they abound, neurotrophins contribute to neuronal growth and survival and to the formation of synaptic pathways. In the mature CNS, however, neurotrophin expression is downregulated, with the exception of brain regions that retain high levels of functional plasticity (Maisonpierre et al. 1990b). It has been suggested that the powerful role played by neurotrophins in development could be harnessed for the treatment of several neurological conditions including injury to the adult spinal cord. This present chapter will first present an overview of the neurotrophin family of neurotrophic factors and their receptors as

well as a brief description of the diverse neurotrophin-mediated signalling pathways that regulate cell survival. This chapter will also examine work that shows that neurotrophins have the capacity to protect spinally projecting neurons against cell death and atrophy as well as to revert the downregulation of regeneration-associated genes that occurs following a spinal cord injury. In this chapter, evidence that the delivery of neurotrophins to the injured spinal cord can outweigh the effect of growth inhibitory molecules present at the injury site will be subsequently reviewed. Moreover, data that demonstrate that the delivery of neurotrophins in the injured cord can promote axonal regeneration through scar tissue will be considered. Finally, this chapter will discuss issues related to recovery of function (or lack thereof) that is reported to accompany axonal regeneration after neurotrophin treatment of the injured spinal cord.

# 2 Members of the Neurotrophin Family of Neurotrophic Factors and Their Receptors

Some of the pro-survival signals that specifically act on neuronal cells belong to a family of target-derived regulatory proteins called neurotrophins, whose main members include nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), neurotrophin-3 (NT-3), and neurotrophin-4 (NT-4). Nerve growth factor (NGF), isolated from snake venom, was the first neurotrophin to be identified (Cohen et al. 1954). Decades later, brain-derived neurotrophic factor (BDNF) was subsequently purified from pig brain tissue (Barde et al. 1982). Years later, neurotrophin-3 (NT-3) was simultaneously characterized by several groups of researchers with the use of molecular cloning techniques (Enfors et al. 1990; Hohn et al. 1990; Rosenthal et al. 1990; Jones and Reichardt 1990; Kaisho et al. 1990; Maisonpierre et al. 1990a). With these techniques, another member of the neurotrophin family, neurotrophin-4/5 (NT-4/5), was identified the following year (Berkemeier et al. 1991; Hallböök et al. 1991). More recently, neurotrophic factor-6 (NT-6) and neurotrophic factor-7 (NT-7), both satisfying the structural criteria that characterize the neurotrophin family, were cloned from a genomic DNA of the platyfish Xiphophorus maculatus and the fish Cyprinus carpio, respectively (Götz et al. 1994; Lai et al. 1998).

Each of these small proteins exhibits a unique molecular profile that determines its particular affinity with one subtype of transmembrane tyrosine kinase (Trk) receptor (recently reviewed in Skaper, 2012a). For instance, TrkA is the specific receptor for NGF (Kaplan et al. 1991; Klein et al. 1991a), while TrkB binds BDNF and NT-4/5 (Klein et al. 1991b; Soppet et al. 1991). NT-3 preferentially interacts with TrkC (Lamballe et al. 1991), but it also bind to TrkA and TrkB although with considerably less efficiency (Glass et al. 1991; Squinto et al. 1991; Klein et al. 1991b; Ip et al. 1993). Thus, the neurotrophins exert their trophic activity through the interaction with transmembrane tyrosine kinase (Trk) receptors to which they bind with distinct as well as with some overlapping degree of specificity (see Ip et al. 1993). The biological action of the different members of the neurotrophin family can also be modulated by the low-affinity nerve growth factor receptor or  $p75^{NTR}$ . The  $p75^{NTR}$  can bind neurotrophins either alone or as a co-receptor that

regulates ligand binding to Trk receptors (Rodriguez-Tébar et al. 1990; Hempstead et al. 1991; Frade and Barde 1998; Bibel et al. 1999).

## 3 Neurotrophin/Trk Receptor Intracellular Signalling

The binding of target-derived molecules of neurotrophins to the extracellular region of two Trk receptor molecules at nerve terminals causes the receptors to associate into active dimers (Ullrich and Schlessinger 1990). In this process, the two Trk dimers are cross-linked through the dimeric ligand but they do not make contact with each other directly (Wehrman et al. 2007). Ligand-induced dimerization of the receptor subsequently leads to phosphorylation of several tyrosine kinase residues within the intracellular domain of the receptor. This process results in the recruitment and activation of a number of downstream signalling effector molecules that initiate the intracellular signalling cascade leading to gene expression and activation (Stephens et al. 1994; Cunningham and Greene 1998).

How the neurotrophin/Trk signal triggered at distal nerve terminals is forwarded to the cell body to elicit neuronal responses has been the focus of much investigation. Although at least two alternative models have been proposed (e.g., Ginty and Segal 2002; Ibáñez 2007), there is mounting evidence in support of the "signalling endosome model" (for a recent review, see Chowdary et al. 2012). This model envisions that the activated receptors are internalized through an endocytosic pathway involving clathrin (Howe et al. 2001; Howe and Mobley 2005; Zheng et al. 2008). These ligand/receptor complexes are then packaged with their downstream signalling molecules into neurotrophic factor receptor-associated protein (NTRAP)mediated specialized membrane-bound vesicles called signalling endosomes (Delcroix et al. 2003; Fu et al. 2010; for related recent articles, see Philippidou et al. 2010; Harrington 2011). These signalling endosomes, which serve as signalling platforms, are subsequently translocated in a retrograde fashion, i.e., from the distal end of axons to the cell soma by dynein-dependent motors along the microtubule tracks (Bhattacharrya et al. 2002; Ha et al. 2008). A similar endosome signalling mechanism has been proposed for the anterograde transport of BDNF from the dendrite to the nucleus (Cohen et al. 2010).

Endosome-bound neurotrophin/Trk receptors complexes mediate several signalling pathways. For instance, they activate the Ras/Erk pathway through the phosphorylation of the cytoplasmic "switch" G-protein Ras that subsequently binds and promotes the translocation of the Raf protein to the plasma membrane (Wennerberg et al. 2005; Reichardt 2006). Raf then activates MEK, which in turn recruits Erk (see Lemmon and Schlessinger 2010, Reichardt 2006 and Huang and Reichardt 2003, for detailed reviews on the Trk-activated signalling pathways). The PI3K/Akt signalling pathway is recruited by the activation of the cytoplasmic "switch" Ras and/or the adaptor protein Gab1 that in turn activates PI3K (Rodriguez-Viciana et al. 1994; Holgado-Madruga et al. 1997). Activated PI3K subsequently activates Akt (Franke et al. 1995). Neurotrophin-dependent Trk activation can also initiate other intracellular signalling cascades such as phospholipase C (PLC- $\gamma$ 1). A downstream effect of the PLC- $\gamma$ 1 pathway is the release of Ca<sup>2+</sup> from reticulum endoplasmic stores, therefore providing a point of convergence between neurotrophin and depolarization-induced cell signalling (Mizoguchi et al. 2002; Blanquet and Lamour 1997; Vaillant et al. 1999). On the other hand, cyclic AMP/protein kinase A (cAMP/ PKA) signalling can also be activated by neurotrophin-recruited Trk receptors.

# 4 Neurotrophins as Cell Survival Regulators

Some of the critical functions of these neurotrophin-mediated signalling pathways are to mediate neuronal survival (Levi-Montalcini 1987; Gosh et al. 1994; Fagan et al. 1996), neurite outgrowth and differentiation (Segal et al. 1995; Barnabé-Heider and Miller 2003), and synaptic plasticity (Canossa et al. 2001; Yamada et al. 2002). A potent role of the PI3K/Akt pathway in neuronal survival (Barnabé-Heider and Miller 2003; Atwal et al. 2000; Vaillant et al. 1999; Holgado-Madruga et al. 1997; Dudek et al. 1997) and in synaptic plasticity (Lin et al. 2001) is well established. The Ras/ERK pathway also contributes to neuronal survival (Impey et al. 1998; Watson et al. 2001), although it seems to be involved mainly in neurogenesis, neuronal growth, and differentiation as well as in axonal elongation (Krishna and Narang 2008; Barnabé-Heider and Miller 2003; Markus et al. 2002; Atwal et al. 2000). The PLC- $\gamma 1$  signalling pathway is not completely understood but there is evidence that it is involved in synaptic plasticity and in the release of neurotrophin-mediated neurotrophin (Canossa et al. 2001; Yamada et al. 2002; Choi et al. 2001; Stephens et al. 1994). As to the cAMP/PKA and Ca<sup>2+</sup>/CaM signalling cascades, there is evidence to suggest that they play a role in neurite extension and outgrowth (Cai et al. 1999; Aglash et al. 2008; Spencer et al. 2008; for a recent review, see Tedeschi 2011).

How does neurotrophin signalling regulate cell survival? The neurotrophinactivated signalling pathways discussed above reach a point of convergence in the cell nucleus where they are potent activators of the nuclear transcription cyclic AMP response element-binding protein (CREB). Indeed, factor neurotrophin-activated Ras/Erk, PI3K/AKT, PLC-7, cAMP/PKA, and Ca<sup>2+</sup>/CaM pathways all mediate the phosphorylation of CREB at Serine 133 (Gonzalez and Montminy 1989; Du and Montminy 1998; Perkinton et al. 2002; reviewed by Lonze and Ginty 2002; Arthur et al. 2004; Spencer et al. 2008). Phosphorylation of CREB subsequently regulates the transcription of the antiapoptotic protein Bcl-2 (Riccio et al. 1999; Bonni et al. 1999; Saini et al. 2004) which, in turn, inhibits apoptosis (reviewed by Yuan and Yankner 2000 and Benn and Woolf 2004). Furthermore, survival is mediated by the activation of Akt, a key component of the PI3K/AKT pathway that leads to the phosphorylation of BAD and Forkhead, two potent effectors that inhibit the proapoptotic pathway (Datta et al. 1997; Brunet et al. 1999). Conversely, activation of the Ras/Erk pathway recruits protein kinase Rsk, the phosphorylation of which also results in the deactivation of the proapoptotic protein Bad (Nebrada and Gavin 1999).

Through their different intracellular signalling pathways, neurotrophins are able to promote cell survival in both a transcription-dependent (e.g., through the recruitment of the transcription factor CREB) and an independent (e.g., through the inhibition of the cell's intrinsic death machinery) manner, therefore shifting the survival/death equilibrium towards survival during ontogenesis as well as throughout adulthood (Finkbeiner et al. 1997).

# 5 Neurotrophins Confer Neuroprotection to Injured Spinal Cord-Projecting Neurons

Corticospinal neurons are located in the sensorimotor cortex and give rise to the corticospinal tract (CST). On the other hand, rubrospinal neurons are found in the red nucleus, a midbrain structure from which the rubrospinal tract (RST) originates. Injury to these long descending fiber tracts results in motor dysfunction below the lesion site. In the adult spinal cord, there is little evidence to suggest that spontaneous regeneration of these motor pathways occurs and that axotomized corticospinal and rubrospinal neurons reinnervate their former target fields (but see Hiebert et al. 2000). What are the response of corticospinal and rubrospinal neurons to axonal transection? It has been long been put forward that the neurons of origin of the rubrospinal tract undergo chromatolysis in reaction to axotomy (Egan et al. 1977). It has also been observed that, in both the sensorimotor cortex and in the red nucleus, the severity of the pathological response is distance specific with more severe pathological reaction, e.g., atrophy and death observed when axonal damage is closer to the cell soma (Egan et al. 1977; Kobayashi et al. 1997).

Giehl and Tetzlaff (1996) showed that axotomy at the level of the internal capsule (i.e., short distance from cell soma) induces death in nearly half the corticospinal neurons, with the surviving neurons displaying severe atrophy (see also Fernandes et al. 1999). However, there is now mounting evidence that the death of axotomized corticospinal neurons can be fully prevented by intraparenchymal delivery of BDNF and NT-3 during the first week after the injury. Indeed, application of BDNF completely alleviates atrophy of corticospinal, whereas the delivery of NT-3 only partially reverses it (Giehl and Tetzlaff 1996). In the same experimental paradigm, the delivery of BDNF, but not NT-3, during the first 2 weeks after the injury promotes the survival of corticospinal neurons for up to 42 days without further exogenous supply of neurotrophins (Hammond et al. 1999). These data suggest that early treatment with BDNF can trigger long-term downstream mechanisms that protect neurons from axotomy-induced death (Hammond et al. 1999).

Infusion of BDNF or NT-4/5 applied on transected rubrospinal fibers (Kobayashi et al. 1997; Bregman et al. 1998) or in the red nucleus (Bretzner et al. 2008) also prevents atrophy of rat rubrospinal neurons. Interestingly, the same group of co-workers have provided evidence that BDNF applied to the cell soma of axotomized rubrospinal neurons promotes their survival up to 1 year after the injury

(Kwon et al. 2002), whereas delayed application of BDNF on truncated rubrospinal fibers (i.e., 2 months after axotomy) failed to reverse atrophy (Kwon et al. 2004). The latter immunohistochemical analysis by Kwon and colleagues revealed that TrkB receptors remain present on the cell bodies of rubrospinal neurons, but not on their axons after axotomy, therefore providing one explanation as to why delayed application of BDNF on truncated rubrospinal axons failed to reverse atrophy (Kwon et al. 2004). The post-axotomy delivery of BDNF, either to the red nucleus or to its transected fibers, has also been reported to reverse neuronal atrophy (Liu et al. 2002; Tobias et al. 2003; Ruitenberg et al. 2004).

# 6 Neurotrophins Upregulate Regeneration-Associated Genes in the Injured Spinal Cord

Axotomy of rubrospinal and corticospinal neurons is accompanied by changes in expression of several genes that are directly involved in regeneration (for a recent review, see Gerin et al. 2011). These include the genes encoding for growthassociated proteins GAP-43 and SCG10, which are known to be involved in the regulation of actin (Frey et al. 2000) and microtubule (Reiderer et al. 1997) dynamics in growth cones during axonal regeneration. A study by Mason and coworkers has demonstrated by means of in situ hybridization that both Gap-43 and SCG10 were upregulated after intracortical lesions that axotomized corticospinal neurons (Mason et al. 2003). Increased GAP-43 mRNA expression in rubrospinal neurons was also reported shortly after axotomy performed at cervical level 3 (C3) (Tetzlaff et al. 1991; Fernandes et al. 1999; Storer and Houle 2003). GAP-43 upregulation is accompanied by an increase in the mRNA encoding both  $T\alpha 1$ and  $\beta$ II-tubulin (Tetzlaff et al. 1991; Storer and Houle 2003), two microtubule proteins that are essential for neurite growth in both developing and regenerating neurons (Hoffman 1989; Gloster et al. 1994; Fausett and Goldman 2006). In the second week after axotomy, however, rubrospinal neurons exhibit a decline in GAP-43 and  $T\alpha$ 1-tubulin expression that coincides with neuronal atrophy (Tetzlaff et al. 1991). Interestingly, the delivery of BDNF and NT-4/5 can maintain the upregulation of GAP-43 and Ta1-tubulin mRNAs expressed by axotomized rubrospinal neurons (Kobayashi et al. 1997; Kwon et al. 2007). The upregulation of GAP-43 and  $\beta$ II- tubulin mRNAs in rubrospinal neurons has also been observed after the application of GDNF, but not BDNF, directly on the truncated axons (Storer et al. 2003). Together, these results show that corticospinal and rubrospinal neurons have the intrinsic capacity to upregulate genes that encode growthassociated proteins in a manner that is similar to that exhibited by neurons capable of regeneration (Hoffman and Cleveland 1988; Miller et al. 1989). These data also suggest that the application of neurotrophins to injured spinally projecting neurons can maintain the axotomy-induced expression of growth-associated genes beyond the second week after the injury.

# 7 Neurotrophins Outweigh the Effects of Inhibitory Molecules in Spinal Cord Injury

The inability of axotomized spinally projecting neurons to regenerate has been attributed, at least partly, to myelin-derived axonal growth inhibitory factors present at the injury site (for reviews, see Silver and Miller 2004; Chaudhry and Filbin 2007; Sharma et al. 2012). Indeed, several myelin-associated inhibitory molecules are able to limit neurite outgrowth and to mediate growth cone collapse. These include Nogo-A (Chen et al. 2000), myelin-associated glycoprotein (MAG) (McKerrecher et al. 1994; Mukhopadhyay et al. 1994), and oligodendrocyte myelin glycoprotein (OMgp) (Mikol and Stefansson 1988; Mikol et al. 1990; recently reviewed in Skaper 2012b). On the other hand, glial scar tissue represents a powerful obstacle to axonal elongation through the lesion, with potent inhibitory extracellular matrix molecules such as chondroitin- and keratan-sulfate proteoglycans produced by reactive astrocytes (Gallo et al. 1987; Gallo and Bertolotto 1990; but see Rolls et al. 2009) and the repellent proteins semaphorin 3 (Pasterkamp et al. 1999: Pasterkamp and Verhaagen 2006; for a recent review, see Pastekamp 2012) and ephrins (Xu et al. 1999; Mellitzer et al. 1999). Since the initial identification of these above-mentioned inhibitory molecules, a wealth of effort has been deployed to reverse the effect of myelin- and scar-derived inhibitory molecule, with strategies such as blocking the effect of Nogo with antibodies (Merkler et al. 2001; GrandPré et al. 2002; Li and Strittmatter 2003; Weinmann et al. 2006; for a review, see Onose et al. 2006), delivering MAG (Vyas et al. 2005) and ephrin (Goldshmit et al. 2004) antagonists and enzymatically digesting chondroitin-sulfate proteoglycans with chondroitinase ABC (Brückner et al. 1999; Starkey et al. 2012; for a recent review, see Bradbury and Carter 2011).

Once thought to be an impenetrable barrier to axonal regeneration, the action of the different inhibitory molecules above mentioned can be counteracted by enhancing the intrinsic growth capacity of axotomized neurons (for recent reviews, see Giger et al. 2010; Hollis and Tuszynski 2011). The regenerative effect of neurotrophins has been extensively studied in rat models of spinal cord injury (for a recent review, see Bo et al. 2011). In the majority of these studies, gene therapy is selected as the method of choice for the delivery of neurotrophins. Briefly, the gene encoding the neurotrophin of choice is cloned into replicationdeficient viral vectors that can be directly delivered at the site of injury (Schnell et al. 1994; Houweling et al. 1998; Namiki et al. 2000; Blits et al. 2003; Koda et al. 2004). In most cases, however, vehicle cells such as fibroblasts (Grill et al., 1997a, b; McTigue et al. 1998; Tobias et al. 2003; Tuszynski et al. 2003; Zhou et al. 2003), permissive autologous bone marrow stromal cells (Koda et al. 2007; Lu et al. 2012), Schwann cells (Xu et al. 1995), neuronal and glial progenitors (Bonner et al. 2010), and olfactory ensheathing glial cells (e.g., Ruitenberg et al. 2003) are modified by viral-mediated transfer of the gene sequence for BDNF. Such virally transduced cells are known to produce biologically active neurotrophins in vitro (e.g., Liu et al. 1999;

Jin et al. 2002; Ruitenberg et al. 2003). These genetically engineered cells are then implanted acutely into the cavity created by the spinal cord injury where they are known to release the therapeutic agent (e.g., Lu et al. 2005). Such BDNF-expressing cell implants have been reported to fill the lesion cavity and to survive in the spinal cord and express the transgene for up to 4 months (Ruitenberg et al. 2003; Tobias et al. 2003).

# 8 Regenerative Property of Neurotrophins on Lesioned Motor Axons

Several groups of researchers have reported that grafts of cells genetically modified to express BDNF create a permissive environment for the injured spinal cord and in particular for the rubrospinal tract. The regenerative properties of engineered cell grafts can be observed even if they are transplanted several weeks after the injury (Tobias et al. 2003). However, the extent to which these BDNF-secreting cell grafts promote axonal growth is somewhat inconsistent. For example, extensive axonal growth has been reported (Lu et al. 2005), sometime extending well outside the boundaries of the BDNF-producing cell implant (Liu et al. 1999, 2002), whereas only modest axonal growth (i.e., within the limits of the implant) has been observed by others (Namiki et al. 2000; Jin et al. 2002; Tobias et al. 2003, 2005; Ruitenberg et al. 2003; Koda et al. 2007). Axons found within the cellular grafts have been shown to be positive for GAP-43 (Tobias et al. 2003, 2005; Koda et al. 2007) and for the marker for axonal neurofilaments TR-97 (Tobias et al. 2003, 2005), therefore providing evidence of motor axonal regeneration. It is worth mentioning that BDNF as well as NT-3-induced axonal regeneration has been associated with oligodendrocyte proliferation and enhanced myelination of ingrowing motor axons (McTigue et al. 1998).

Nearly two decades ago, Schnell and co-workers have reported that injections of NT-3, but not BDNF, result in modest regenerative sprouting of the CST (Schnell et al. 1994). Similar results were obtained a few years later with NT-3 dissolved into a collagen matrix that was implanted in the lesioned spinal cord (Houweling et al. 1998). As is the case for BDNF, viral-mediated gene therapy involving NT-3 as the transgene has been widely used in acute and chronic animal models of spinal cord injury. Grill and co-workers (1997b) implanted primary rat fibroblasts genetically modified ex vivo to express NT-3 in dorsal spinal cord hemisections that include the CST. Three months later, significant CST growth was reported at, as well as for up to 8 mm distal to, the lesion site, showing the therapeutic benefits of NT-3 in this descending motor fiber pathway (Grill et al. 1997b). Grafts of NT-3-expressing fibroblasts in the injured cord also induce proliferation of oligodendrocytes and encourage myelinogenesis (McTigue et al. 1998). In a study performed by Tuszynsky and co-workers (2003), primary fibroblasts genetically modified to express human NT-3 were implanted in and distal to the lesion cavity after chronic

spinal cord injury. With this dual implantation protocol (e.g., in and distal to the lesion), the elongation and arborization of CST axons was observed up to 15 mm distal to the injury site. Multiple combinational therapy, with cAMP injections into brainstem reticular motor neuron area to encourage their growth state, implants of bone marrow stromal cells to provide a cellular bridge through the spinal cord lesions, and the delivery of BDNF-expressing viral vectors caudal to the lesion sites to attract axonal growth, has recently been performed (Lu et al. 2012). This complex combinational treatment leads to motor axon regeneration through and beyond the limits of the implant, with some evidence of synapse formation with BDNF-expressing neurons caudal to the implant.

Retrograde neural tracers are known to be internalized by nerve endings and transported in a retrograde fashion to the corresponding spinal cord motor neurons (Tosolini and Morris 2012). The retrograde machinery of neurons can also be used to deliver different transgenes to spinal cord motor neurons (Morris et al. 2004). Intramuscular injections of viral vectors containing the gene sequence for diverse neurotrophins have been used as an alternative method to cell-mediated delivery of neurotrophic genes in the spinal cord (Ghage et al. 1995; Baumgartner and Shine 1998a, b; Nakajima et al. 2007; Uchida et al. 2008, 2012; Zhang et al. 2010). Interestingly, it is not necessary to target all motor neurons to achieve sustainable levels of chemoattractant, as non-transduced motor neurons in the vicinity have been shown to internalize neurotrophins through a paracrine mechanism (Baumgartner and Shine 1997). This "bystander" effect greatly enhances the efficacy of neurotrophin gene interventions that take advantage of the retrograde machinery of neurons (e.g., Aharoni et al. 2005). Virally mediated retrograde delivery of the gene sequence for BDNF has been reported to protect motor neurons from apoptosis after a spinal cord injury in rats (Nakajima et al. 2007, 2010) and mice (Uchida et al. 2008, 2012). However, although the delivery and expression of BDNF in motor neurons near a spinal cord lesion should presumably encourage axonal regeneration, no evidence supporting this assumption is currently available. On the other hand, adenoviral vector carrying the gene sequence for NT-3 has been delivered to level motor neurons at lumbar levels via injections in the sciatic nerve in a rat model of unilateral CST lesion at the level of the hindbrain (Zhou et al. 2003). In this study, sprouting from the intact CST (i.e., contralateral to the injection of NT-3-expressing viral vector) was reported to cross the midline at lumbar levels L3/6, suggesting that this particular observation is an instance of NT-3-mediated axonal plasticity rather than true regeneration. The effect of implant-mediated transgenic delivery of NGF, GDNF, and NT-4/5 has been less investigated. However, cellular implants secreting NGF (Grill et al. 1997a; McTigue et al. 1998; Jones et al. 2003), GDNF (Blesch and Tuszynski 2003), and NT-4/5 (Blesch et al. 2004) have been shown to produce significant regeneration of specific motor pathways through the implants. Taken together, these data provide evidence that the delivery of growth-promoting agents such as neurotrophins, if available in sufficient concentration, can outweigh the effect of inhibitory molecules present in the scar tissue.

# 9 Regenerative Properties of Neurotrophins for Lesioned Sensory Axons

Rhizotomy, or crush lesions of the dorsal roots, has been used extensively to study sensory axon regeneration within the spinal cord (for a recent review, see Smith et al. 2012). In these injury models, centrally projecting sensory axons regenerate rapidly until they reach the transition between the peripheral and central nervous system, an area called the dorsal root entry zone (DREZ). Several neurotrophin therapies have been conducted in an effort to encourage sensory axons to reenter the spinal cord after dorsal root crush. After such lesions, intraspinal infusion of NGF (Odega and Hagg 1996; Ramer et al. 2000), NT-3 (Zhang et al. 1998; Ramer et al. 2000, 2001), and GDNF, but not BDNF (Ramer et al. 2000), has been shown to promote extensive regeneration of sensory axons through the DREZ and in the spinal cord. Dorsal horn injections of adenovirus carrying the gene sequence for NGF have been reported to induce robust regeneration of dorsal root sensory axons through the DREZ and into former target laminae II and IV in the dorsal horn of the cord (Romero et al. 2001). In this study, however, sensory afferent axons were also observed to innervate laminae III and VI in the dorsal horn as well as ventral horn laminae VII-X. In a follow-up study, the same group of researchers was able to confine the regenerating sensory axons to the superficial layers of the dorsal horn by coupling the injection of the NGF-expressing vector with that of a vector encoding the gene for the negative guidance molecule semaphorin 3A in the ventral horn (Tang et al. 2007).

Another model to study regeneration of sensory axons is to create a partial spinal cord lesion that disrupts the ascending dorsal column. With this model, Taylor and colleagues (2006) have shown that a continuous gradient of lentivirus-mediated neurotrophin-3 (NT-3) in and immediately rostral to a lesion elicit modest axonal growth beyond the lesion. However, the sensory axons were found to stop elongating when they reach the rostral limit of the NT-3 gradient (Taylor et al. 2006). The same group of researchers has further demonstrated that axons have the potential to reach and reconnect with their former target if the distance between the lesion and the target is very small (2 mm) and is filled with a continuous gradient of NT-3 (Taylor Alto et al. 2009). Together, these findings show that damaged axons in the adult animal not only have the potential to elongate but also can form new synapses with former targets and that neurotrophins play a critical role in this process.

# 10 Functional Recovery After Neurotrophin Treatment of the Injured Spinal Cord

In the previous sections of this chapter, evidence that severed motor and sensory fiber pathways can regenerate through scar tissue, if neurotrophins are made available, has been reviewed. Is neurotrophin-mediated axonal regeneration associated with the recovery of function? Unfortunately, the answer is not straightforward. Of the various experimental models of spinal cord injury, dorsal rhizotomy seems to be particularly amenable to recovery of function after neurotrophin treatment. In particular, the delivery of NGF after dorsal rhizotomy at lumbar levels 4 and 5 (L4/5) has been associated with near complete recovery of thermal nociception (Romero et al. 2001; Tang et al. 2007). Particularly, the report by Ramer et al. (2000) that deafferented dorsal horn sensory neurons can be synaptically driven by peripheral nerve stimulation after NGF, NT-3, and GDNF treatment is strong evidence of functional recovery. A therapeutic effect for BDNF on the recovery of thermal sensitivity has also been reported (Shumsky et al. 2003) (see Table 1).

Several teams of scientists have provided evidence that BDNF treatment after cervical lesions that involve the RST is accompanied by various degrees of recovery of function. After such lesions, BDNF delivery has been reported by different teams of researchers to enhance performance on the rope walking (Kim et al. 2001; Ruitenberg et al. 2003; Tobias et al. 2005) and cylinder tests (Liu et al. 1999; Tobias et al. 2005). Open-field testing with the BBB locomotor rating scale (Basso et al. 1995), on the other hand, has led to conflicting results. Indeed, significant improvement on the BBB open-field locomotor test has been reported in BDNF-treated animals (Koda et al. 2004), whereas complete recovery on this test has been shown to be independently of the nature of the treatment, i.e., fibroblasts expressing BDNF or control fibroblasts (Kim et al. 2001; Tobias et al. 2005). Together, these findings therefore suggest that BBB score is not useful to assess recovery of motor function after cervical lesions that target the RST. It is worthwhile to point out that the rope walking and the narrow beam tests as well as the BBB open-field locomotor test assess motor performance involving whole limb movement per se, or in relation to the intact limbs. However, evidence that lesions restricted to the RST do not lead to deficits in whole limb movement (Whishaw et al. 1998; Morris et al. 2011) suggests that these above-mentioned behavioral tasks are not particularly stringent to evaluate functional recovery of the RST. Moreover, the return of performance to, or near, preoperative values on some of these tests may therefore reflect compensation of function mediated by other motor pathways, particularly when recovery of function is not accompanied by evidence of RST growth in and through the BDNF-secreting implant.

Significant recovery on the gridwalk test has been reported after dorsal thoracic hemisections that include the main component of the CST, followed by treatment with NT-3-expressing cells (Grill et al. 1997b; Houweling et al. 1998). In both studies, amelioration of locomotor capacity on this test was accompanied by growth of CST axons in the cellular implants. However, the report by Grill and co-workers (1997) that smaller lesions that still abolish the CST are not sufficient to create a deficit on the gridwalk test (Grill et al. 1997b) indicates that the observed CST growth and recovery on the gridwalk test are two independent events. Furthermore, this observation also suggests that multiple pathways, possibly including the lateral aspect of the CST and the RST, contribute to locomotor function as measured by the gridwalk test.

Authors	Type of injury	Treatment	Anatomical outcome	Behavioral testing	Functional recovery after treatment
Tang et al. (2007)	Dorsal rhizotomy at L4/5	Ad-mediated NGF gene and semaphorin3A transfer in dorsal and ventral horns, respectively	Sensory axon growth across DREZ and into superficial dorsal horn laminae I and II	Response to noxious thermal stimulation	Functional restoration of nociception
Ramer et al. (2000)	Dorsal rhizotomy between C4 and T2	Delivery of NGF, NT-3, BDNF, and GDNF	Sensory axon growth across DREZ and into the spinal cord after treatment with NGF, NT- 3, GDNF but not BDNF	Peripheral nerve stimulation	Dorsal horn neurons found to be synaptically driven with NGF, NT-3, GDNF but not BDNF
Romero et al. (2001)	Dorsal rhizotomy at L4/5	Delayed Ad- mediated NGF gene transfer in dorsal horn	Robust axonal growth in normal and aberrant locations with the dorsal horn	Plantar heat test and grid runway	Near complete recovery of hindlimb thermal nociception
Shumsky et al. (2003)	Unilateral hemisection at C3/4	Delayed delivery of BDNF and NT- 3-expressing fibroblasts	Not reported	Battery of motor and sensorimotor tasks	No recovery on motor tasks but modest recovery of thermal sensitivity
Ruitenberg et al. (2003)	Unilateral DLF lesion involving the RST at C4	Acute delivery of BDNF- expressing OECs	Sprouting of RST axons in graft	Rope walking	Improved hindlimb performance
Liu et al. (1999)	Unilateral LF lesion involving the RST at C3/4	Acute delivery of BDNF- expressing fibroblasts	Growth of RST axons 3–4 segments below grafts	Cylinder test	Significant recovery
Tobias et al. (2005)	Unilateral DLF lesion involving the RST at C4	Acute delivery of BDNF- expressing fibroblasts	No evidence of growth of RST fibers in the implant	Rope walking test, cylinder test, BBB locomotor test	Complete recovery on BBB in treated and control rats, partial recovery in treated rats on cylinder test, partial recovery of rope walking in treated and control rats

**Table 1** Compilation of several scientific reports that have addressed recovery of function after diverse neurotrophin treatments

(continued)

	,				
Authors Koda et al.	Type of injury Complete	Treatment Ad-mediated	Anatomical outcome Growth of RST	Behavioral testing BBB	Functional recovery after treatment Significant
(2004)	transection at T8	BDNF gene transfer in both stumps	axons through the lesion site	locomotor test (Basso et al. 1995)	improvement for at least 6 weeks
Kim et al. (2001)	Unilateral LF lesion at C3/4	Acute delivery of BDNF- expressing fibroblasts	Not reported	Swim test, patch-removal test, rope walking test, BBB locomotor test, narrow beam test,	No recovery on swim test, shorter latency to touch the patch, significant recovery on the recovery on BBB locomotor, and narrow beam tests in treated and control rats
Blits et al. (2003)	T9/10 excision	Schwann cell bridge and acute delivery of AAV- mediated BDNF and NT- 3 gene transfer 5 mm caudal to bridge	Axonal growth in the cellular bridge but not beyond its caudal limits	BBB locomotor scale	Modest improvement of hindlimb function
Namiki et al. (2000)	Compression injury at T3	Infusion of BDNF and NT- 3 for 14 days after injury	Smaller lesion size and regeneration through the injury	Inclined plane technique	Significant recovery at 1 week after injury. No significant recovery at later times
Houweling et al. (1998)	Dorsal hemisection between T8 and T10	NT-3 in a collagen matrix applied at the site of the injury	containing collagen matrix		Significant recovery
Grill et al. (1997b)	Dorsal hemisection at T7	Acute delivery of NT-3- expressing fibroblasts	Significant growth of CST axons in cellular implant	Gridwalk test	Significant recovery
					(continued)

## Table 1 (continued)

(continued)

Authors	Type of injury	Treatment	Anatomical outcome	Behavioral testing	Functional recovery after treatment
Tuszynski et al. (2003)	Dorsal hemisection at T7	Delayed delivery of NT- 3-expressing fibroblasts placed in, and caudal to the injury site	Significant growth of CST axons up to 15 mm distal to the injury site	BBB locomotor scale	Modest recovery

Table 1 (continued	nued)	(0	1	le	ab	T
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In an effort to encourage axonal growth below the cell implants, Blits et al. (2003) used Schwann cell implants to bridge thoracic spinal cord transections and subsequently delivered adeno-associated viral vectors containing the gene sequence for BDNF and NT-3 caudal to the cellular bridge. This surgical treatment, however, failed to promote growth caudal to the level of the NT-3-secreting bridge. Using a similar strategy, Tuszynski and co-workers (2003) observed growth of CST axons up to 15 mm distal to the injury level after the implantation of NT-3-expressing fibroblasts in, as well as caudal to a thoracic dorsal hemisection. Despite the considerable difference in the extent of axonal growth reported in these two studies, both procedures led to equally modest recovery on the BBB locomotor scale (Blits et al. 2003; Tuszynski et al. 2003). Thus, the striking difference in the extents of NT-3-mediated regeneration observed between these studies is by no means reflected in the level of performance on the BBB locomotor scale. In summary, that axonal regeneration after neurotrophin treatment has a direct impact on the recovery of function, at least with regard to the motor modality, is yet to be established.

### 11 Conclusion

In its infancy only a few decades ago, spinal cord regeneration has become a vast and dynamic field of research in neuroscience. Over the last 30 years, many groups of scientists have concentrated their efforts to identify growth inhibitors that are present in the damaged spinal cord and to develop strategies to counteract their action. Others have sought to design ways to increase the levels of growthpromoting neurotrophin expression at the site of injury in an attempt to offset the effect of these inhibitory molecules. There is now compelling evidence that neurotrophic factors can promote the survival of neurons giving rise to the different ascending and descending fiber pathways running in the spinal cord and that are axotomized after an injury. Moreover, it is well substantiated that the action of inhibitory signalling molecules that act as a barrier to axonal regeneration can be reverted by the delivery of neurotrophins to the injured spinal cord. Collectively, these key findings represent significant advances towards the development of therapeutic strategies to promote axonal regeneration through a spinal cord lesion. Further work such as subjecting promising anatomical outcomes to independent replication is required before these preclinical data can be translated to clinical trials.

Recovery of function after spinal cord injury remains an enormous challenge for neuroscientists working in the field of spinal cord regeneration. Indeed, although the literature examined in the present chapter has reported axonal growth in and even several millimeters below the level of injury, there is no compelling evidence to show that the regenerative benefits of neurotrophin treatment are directly associated with functional recovery. Several factors may contribute to this unfulfilled endeavor. First and foremost, functional recovery must be supported by anatomical evidence of reinnervation of former postsynaptic target cells. This includes the demonstration that appropriate synaptic contacts (i.e., axodendritic synapses) with the postsynaptic target cells have been reestablished (e.g., Taylor Alto et al. 2009). Second, behavioral evidence for functional recovery gains significant strength if supported by electrophysiological data. In this regard, the report by Ramer et al. (2000) that deafferented sensory neurons in the dorsal horn can be synaptically driven by stimulation of the peripheral nerve after neurotrophin treatment constitutes powerful evidence of functional recovery. The years to come will certainly bring us closer to resolving this difficult task.

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# **Neurotoxicity and Stroke**

# Victoria O'Collins, David Howells, and Romesh Markus

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#### Abstract

Stroke is the clinical manifestation of the occlusion (ischemic stroke) or rupture of a blood vessel to the brain (hemorrhagic stroke). Neurotoxicity is triggered in stroke by the failure of oxygen delivery and the buildup of metabolites resulting in a cascade of harmful physiological events. Here, we review and contrast neurotoxicity and its corollary – neuroprotection – in the context of preclinical

V. O'Collins • D. Howells

R. Markus (🖂)

Florey Neuroscience Institutes, Melbourne Brain Centre, Heidelberg, VIC, Australia e-mail: ocollins@unimelb.edu.au; david.howells@unimelb.edu.au

Department of Neurology, St. Vincent's Hospital, Darlinghurst, Sydney, NSW, Australia e-mail: rmarkus@stvincents.com.au

(animal) and clinical (patient) drug trials. Preclinical tests are distinguished by greater homogeneity, earlier time windows for treatment, and greater probability of success. Many of the same processes have been targeted in preclinical and clinical trials – such as excitotoxicity, oxidation, inflammation, and trophic factors – however, it is not a seamless flow from bench to bedside and back. Current trends in stroke neuroprotection include combination neuroprotection and improved trial designs. Areas where drug development in stroke could be improved include better target identification, development of lead drug candidates prior to preclinical testing and use of computational tool for prediction of toxicity based on biochemical pathway and drug characteristics.

#### Keywords

Neurotoxicity • Stroke • Neuroprotection • Brain ischemia • Animal models • Clinical trials

#### List of Abbreviations

AMPA	3-hydroxy-5-methyl-4-isoxazole propionic acid (AMPA)
AO	Antioxidant
FRS	Free radical scavenger
ICA	Internal carotid artery
MCA	Middle cerebral artery
MCAo	Middle cerebral artery occlusion
NMDA	N-methyl-D-aspartate
tPA	Tissue plasminogen activator
WHO	World Health Organization

# 1 Introduction to Stroke

# 1.1 Definition of Stroke

Stroke is the clinical manifestation of the occlusion (ischemic stroke) or rupture of a blood vessel to the brain (hemorrhagic stroke). It has been formally defined by the World Health Organization as "rapidly developing clinical signs of focal (or global) disturbance of cerebral function lasting more than 24 hours (unless interrupted by surgery or death) with no apparent cause other than a vascular origin" (WHO 1988). This review is limited to ischemic stroke which accounts for approximately 85 % of strokes in humans.

# 1.2 Pathophysiology of Ischemic Stroke

Knowledge of the mechanisms by which ischemic stroke causes brain damage has been derived from autopsy, from animal experiments, and more recently from

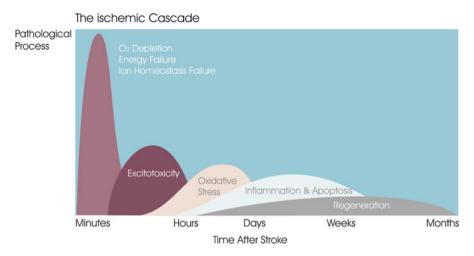


Fig. 1 The ischemic cascade (Adapted from Heiss et al. 1999)

imaging of the human brain. These data suggest that the extent to which blood flow is reduced in the brain is an important determinant of the response of cells (Astrup et al. 1981; Bandera et al. 2006).

Occlusion of a cerebral artery results in a reduction in oxygen, glucose, and nutrients in the territory of the brain supplied by the artery. The brain attempts to compensate for small reductions in blood flow (hypoperfusion) by recruiting blood flow through collateral vessels and by autoregulatory vasodilation (Derdeyn et al. 1999). Further reductions in blood flow are compensated for by increasing the amount of oxygen extracted from blood (Derdeyn et al. 1999). A failure to normalize perfusion pressure and blood flow will result in cellular swelling, disturbances in the redox state, lactic acidosis, loss of membrane dynamics, and ultimately cell death (Derdeyn et al. 1999).

On the basis of the degree to which blood flow and metabolism are compromised, brain tissue is often divided into normal healthy tissue, oligemic tissue, penumbral tissue, and infarcted tissue, in order of increasing severity of damage. Infarction initially occurs in the most severely hypoperfused region ("core"). The surrounding nonfunctional but potentially salvageable brain tissue ("penumbra") progresses to infarction with time as a result of a cascade of ischemia-induced biochemical changes resulting in cellular death (Markus et al. 2003).

Time is a critical determinant of the viability of brain tissue following a stroke. Evidence from animal models of stroke has demonstrated that the occlusion of a blood vessel sets in train a series of physiological events which have been termed the "ischemic cascade" (Dirnagl et al. 1999; Heiss et al. 1999). Primary damage following the loss of blood flow includes oxygen depletion, energy failure, and the failure of ion homeostasis (Fig. 1) (Heiss et al. 1999).

#### 1.3 Neurotoxicity and Stroke

Ischemic stroke is a complex, multifaceted, and heterogeneous disease; thus, neurotoxicity is not easy to define in the context of stroke. Whilst the lack of oxygen and nutrients is the ultimate "neurotoxin," secondary damage may result from the following mechanisms: (1) "excitotoxicity," a surge in extracellular excitatory amino acids such as glutamate which become toxic because they are not buffered by normal cellular reuptake mechanisms (mainly astrocytes) (Benveniste et al. 1984; Benveniste 1991; Rothman and Olney 1986); (2) "oxidative stress," the massive release of free radicals that follow when endogenous production mechanisms become overactive (e.g., NOX enzymes) or when protective mechanisms (e.g., glutathione, superoxide dismutase) are swamped (Uenohara et al. 1988); and (3) inflammation and associated cell death, the inflammatory and apoptotic response marked by the migration and activation of microglia and macrophages at the lesion site and apoptotic signaling mechanisms (e.g., misplaced mitochondrial proteins which lead to further cell death) (Abraham and Lazar 2000; Mabuchi et al. 2000). Neurotoxicity may also result from disturbances in physiological processes such as acidosis, hyperthermia, blood sugar imbalance, and excessively high or low blood pressure.

Neurotoxicity may also be used to describe side effects of medications administered to manage stroke or a patient's comorbidities. For instance, tissue plasminogen activator (tPA) – expressed endogenously and administered in recombinant form therapeutically to break down clots – is a lifesaving therapy, but in animal models it has also been associated with neuronal damage, especially if the blood– brain barrier is not intact or if administered outside the vascular compartment (Wang et al. 1998, 1999; Kaur et al. 2004). Similarly, early agents targeting excitotoxicity (e.g., NMDA antagonists like MK-801) had poor profiles in terms of their effect on cognition (Muir and Lees 1995).

Neurotoxicity, however, is not the most common term designating damage in stroke research; when it does, it is often when implicating substances related to damage in other areas of medicine such as iron-induced neurotoxicity (Selim and Ratan 2004). Blood is neurotoxic if it gets outside of the vessels; the iron in hemoglobin is toxic to the brain and blood-signaling proteins like cytokines and chemokines. This causes inappropriate stimulation of the neuraxis and plays a role in directing invasion of inflammatory cells which themselves cause further damage but also help in regeneration. These regenerative responses are believed to be activated in the days and weeks following stroke onset (Wiltrout et al. 2007), but by this time most of the damage has been done.

One reason why endogenous substances upregulated during the ischemic cascade are not automatically designated as neurotoxic is that they might also take on neuroprotective roles, depending upon the time after iachemic onset and cell type. For instance, microglia has both neurotrophic and proinflammatory roles (Kim and de Vellis 2005) and may play a complex role after ischemia. Neuroprotection, rather than neurotoxicity, has therefore been the focus of intense interest in stroke research over the past 25 years.

# 2 Neuroprotection and Stroke

### 2.1 Clinical Management of Acute Stroke

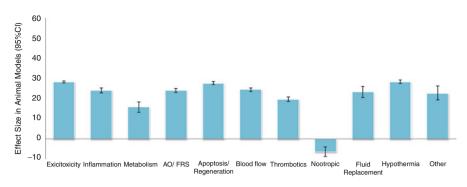
Current evidence supports the use of a limited number of acute stroke interventions (Donnan et al. 2008): (1) management of patients in hospital stroke units, (2) reestablishment of blood flow using recombinant tissue activator (tPA) (The National Institute of Neurological Disorders and Stroke rt-PA Stroke Study Group 1995), (3) aspirin (The National Institute of Neurological Disorders and Stroke rt-PA Stroke Study Group 1997; International Stroke Trial Collaborative Group 1997), and (4) hemicraniectomy (Vahedi et al. 2007). Of these treatments, the most significant and widely available gains during the acute phase of stroke have been secured through caring for patients in a dedicated hospital stroke unit (Donnan et al. 2008). The benefits of these interventions are not mutually exclusive; however, for patients in hospital stroke units are probably more likely to have access to the other three interventions.

Therapies that target the cerebral vasculature by restoration of cerebral blood flow (thrombolytics, mechanical embolectomy) and preventing clot formation (aspirin) have shown unequivocal benefit in clinical trials and are approved for use in clinical practice. Neuroprotective strategies targeting the injurious biochemical and molecular cascade are designed to restrict injury to the ischemic brain by preventing neuronal cell death, especially in the salvageable penumbral region. While the concept of neuroprotection has shown promise in experimental studies, thus far there has been failure of translation into clinical success.

# 2.2 Neuroprotection and Animal Models of Stroke

Over the past 50 years, animal models have been used to test the efficacy of potential neuroprotective stroke therapies both before and after they have gone to clinical trial. Most commonly, these animal models entail occlusion of the middle cerebral artery to induce a "focal" area of brain damage. The standard method of evaluating treatments has been by applying a dye or stain to fine slices of brain tissue after the animal has been sacrificed. Various functional tests for animals have been derived from their clinical counterparts and purport to measure sensory and motor deficits. Such tests have been the mainstay of preclinical testing of potential therapies to reduce the impact of stroke.

To obtain global estimates of efficacy in animal models, data were pooled from two systematic reviews (O'Collins et al. 2006, 2012), which identified studies reporting the efficacy of interventions in animal experiments published 1978–2010. Testing the effect of therapies on infarct size was undertaken in 3,288 acute ischemic stroke experiments (47,899 animals); 94 % of these experiments were undertaken in rats or mice. Data were combined using meta-analysis and meta-regression, partitioned on the basis of stroke model



**Fig. 2** Testing of acute stroke treatments in animal models. *Bar chart* of data from 3,288 controlled experimental tests of therapies in acute stroke in animal models (47,899 animals), subject to meta-analysis. A higher effect size indicates greater efficacy of the therapeutic class, compared to the control condition. Abbreviations: *AO* antioxidant, *FRS* free radical scavenger

and therapeutic target (e.g., excitotoxicity, thrombolysis, oxidative stress). Because of the heterogeneity, we used random effects meta-analyses (DerSimonian and Laird 1986).

Across all 502 potential therapies included in the dataset, the studies indicated on average a substantial reduction in brain damage (infarct) through use of a therapy (effect size =  $25.1 \pm .15$ , 95%CI 24–25.4). Of the 3,288 experiments, 24 % involved testing of excitotoxicity agents, 17 % involved agents targeting apoptosis or regeneration, 13 % involved free radical scavengers/antioxidants, and 12 % deployed hypothermia. The most effective of the therapeutic classes was hypothermia (effect size =  $28.4 \pm 0.4$ , 95%CI 27.6–29.3) and anti-excitotoxic drugs ( $28.3 \pm 0.3$ , 95%CI 27.7–28.8) (see Fig. 2). However, classification of treatment mechanisms is notoriously difficult with treatments with multiple mechanisms of action and no objective measure of the strength of action in a domain.

# 2.3 Clinical Trials of Neuroprotection in Stroke Versus Animal Data

The Internet Stroke Center website provides a comprehensive database of neuroprotective agents that have been studied in humans (Internet Stroke Center 2013). Table 1 summarizes the key Phase III neuroprotection trials in humans, none of which have shown unequivocal clinical benefit (Table 1). Details of the drugs tested are given below. No neuroprotective strategy – in a narrow sense of the term – is currently approved for clinical use. By contrast, individual therapies tested in animal models and taken to Phase III clinical trial, estimates of efficacy from meta-analysis (Table 2) tend to suggest a positive effect. Note that these estimates are representative and not exhaustive of all animal model studies for each of the drugs listed below.

Category	Drug name	Mechanism	Time window	Time Patients Clinic window (drug/placebo) Phase	Clinical Phase	Year Trial	Trial	Reference
Antioxidant	NXY-059	Free radical scavenger	9	1,646/1,660	Ш	2007	SAINT II	(Shuaib et al. 2007)
	NXY-059	Free radical scavenger	6	853/852	Ш	2006	2006 SAINT I	(Lees et al. 2006)
	Tirilazad	Free radical scavenger	6	276/280	Ш	1996	1996 RANTTAS	(The RANTTAS
								Investigators 1996)
Calcium	Nimodipine	Calcium antagonist	9	225/229	III	2001	2001 VENUS	(Horn et al. 2001)
modulators	Nimodipine	Calcium antagonist	48	800/264	III	1992		(The American Nimodipine
								Study Group 1992)
	Nimodipine	Calcium antagonist	48	607/608	III	1990	1990 TRUST	(Trust Study Group 1990)
	Ginsenoside	Calcium antagonist	72	290/96	Ш	2012		(Liu et al. 2012)
Excitotoxicity Aptiganel	Aptiganel	NMDA ion channel blocker	12	414/214	III/II	2002		(Albers et al. 2001)
	Cervene	Opioid antagonist	9	163/167	Π	2000		(Clark et al. 2000)
	(Nalmefene)							
	Clomethiazole	GABA agonist	12	599/599	III	2002	2002 CLASS-1	(Lyden et al. 2002)
	Clomethiazole	GABA agonist	12	680/680	III	1999	CLASS	(Wahlgren et al. 1999)
	Diazepam	GABA agonist	12	426/417	III	2006	2006 EGASIS	(Lodder et al. 2006)
	Gavestinel (GV150526A)	NMDA glycine antagonist	9	701/666	Ш	2001	2001 GAIN AMERICAS (Sacco et al. 2001)	(Sacco et al. 2001)
	Gavestinel (GV150526A)	NMDA glycine antagonist	9	891/897	Π	2000	2000 GAIN INTERNATIONAL	(Lees et al. 2000)
	Lubeluzole	Sodium/calcium channel blocker/NOS inhibitor	6	365/360	Π	1998		(Diener 1998)

 Table 1
 Neuroprotection in acute ischemic stroke. Phase III trials in humans

			Time	Patients	Clinical		
Category	Drug name	Mechanism	window	window (drug/placebo) Phase	Phase	Year Trial	Reference
	Magnesium	NMDA antagonist/Ca channel blocker	12	11,292/1,297	Π	2004 IMAGES	(Muir et al. 2004)
	Selfotel (CGS-19755)	NMDA antagonist	9	281/286	Ш	2000	(Davis et al. 2000)
Membrane	Piracetam	AMPA (Na) modulator	12	464/463	Ш	1997 PASS	(De Deyn et al. 1997)
stabilizers	Citicoline	Phospholipid synthesis	24	1,148/1,150	Ш	2012 ICTUS	(Dávalos et al. 2012)
	Repinotan	Serotonin antagonist	4.5	342/337	III	2009	(Teal et al. 2009)
Trophic	Cerebrolysin N	Neurotrophic	12	529/541	III	2012 CASTA	(Heiss et al. 2012)
factor	Erythropoietin	Anti-apoptosis, oxygen delivery	9	266/256	III	2009	(Ehrenreich et al. 2009)
	Trafermin	Growth factor	6	286	III/II	2002	(Bogousslavsky et al. 2002)

Table 1 (continued)

Category	Drug name	Mechanism	Time window	Number of animals (drug/ control)	No. Experiments	Effect size 土 SE	95%CI	Year
Antioxidant	NXY-059	FRS	5 min	217/117	25	$38 \pm 2$	[33,42]	1999
	Tirilazad	FRS	10 min	205/135	23	$26 \pm 2$	[23,35]	1991
Calcium	Ginsenoside	Calcium antagonist	0 min	217/79	29	$41 \pm 3$	[36,47]	2004
modulators	Nimodipine	Calcium antagonist	-5 min	306/217	36	$24 \pm 2$	[21,27]	1986
Excitotoxicity	Aptigane1 (Cerestat)	NMDA ion channel blocker	15 min	100/52	10	$68 \pm 3$	[63,73]	1993
	Cervene (Nalmefene)	Opioid antagonist	I	0/0	0			None
	Clomethiazole	GABA agonist	15 min	60/39	8	$47 \pm 3$	[41,54]	1995
	Diazepam (Valium)	GABA agonist	30 min	5/5	1	$-13 \pm 25$	[-61, 35]	2004
	Gavestinel (GV150526A)	NMDA glycine antagonist	7 min	161/70	21	$65 \pm 8$	[49,80]	1997
	Lubeluzole	Sodium/calcium channel blocker/ NOS inhibitor	1 min	231/115	19	$40 \pm 2$	[37,43]	1996
	Magnesium sulfate	NMDA antagonist/Ca channel blocker	-12.5 min 125/71	125/71	14	$25 \pm 3$	[19,307]	1996
	Selfotel (CGS- 19755)	NMDA antagonist	5 min	31/21	5	$33 \pm 2$	[28,37]	1990
	YM872	AMPA receptor antagonist	10 min	180/112	18	$24 \pm 3$	[30,98]	1999
Membrane	Citicoline	Nootropic	1 h	481/185	48	$27 \pm 1$	[24,29]	1996
stabilizers	Piracetam	AMPA (Na) modulator	6 h	25/22	4	$10\pm 5$	[0, 20]	2002
Nootropic/ stimulant	Cerebrolysin	Nootropic	24 h	120/40	12	8 土 4	[0,17]	2007
<b>Trophic factors</b>	Erythropoietin	Anti-apoptosis oxygen delivery	6 h	157/65	20	$19 \pm 2$	[15, 23]	1999
	Trafermin <sup>a</sup>	Growth factor	30 min	248/174	29	$32 \pm 2$	[28, 36]	1994
Others	Repinotan	Serotonin antagonist	1 min	192/108	19	$53\pm 1$	[50, 54]	1998

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Neurotoxicity and Stroke

#### 2.3.1 Glutamate Antagonists

Glutamate excitotoxicity is an important mediator of ischemia-induced neural injury via the *N*-methyl-D-aspartate (NMDA) and the 3-hydroxy-5-methyl-4-isoxazole propionic acid (AMPA) receptors.

#### **NMDA Receptor Antagonists**

Aptiganel (CNS-1102), a noncompetitive NMDA receptor antagonist (Albers et al. 2001), and Selfotel (CGS 19755) (Davis et al. 2000), a competitive NMDA receptor antagonist, have been studied in Phase III clinical studies. Although preclinical studies indicated that neuroprotection with these agents occurred only after very early administration, a time window of up to 6 h after stroke onset was utilized in human studies. Both agents exhibited significant neurobehavioral toxicity, resulting in a lower dosage being used in human studies than that predicted to provide neuroprotection from comparative preclinical animal studies. A higher mortality was observed in the Aptiganel group with a trend towards higher mortality in the Selfotel group, raising the possibility that NMDA receptor-mediated neuronal survival mechanisms were adversely affected by the delayed administration (Hoyte et al. 2004).

NMDA receptor antagonists have been relatively well studied in animal models of stroke compared with other classes, although even amongst these drugs, relatively few tests were published prior to clinical trial. Aptiganel has been studied in rat models of stroke and has been found to reduce infarct size when administered minutes after stroke onset with effects studied within the few days (Minematsu et al. 1993; Pitsikas et al. 2001; Zhao et al. 2010). When administered subcutaneously in Fisher 344 rats – as opposed to intravenously in more commonly used rat strains – beneficial effects were only seen in a narrow dose range (Pitsikas et al. 2001).

Selfotel has been investigated in focal ischemia models, global ischemia, and spinal cord ischemia, both in cat focal ischemia models (Okada et al. 1997) and in rat models (Jolkkonen et al. 1999; Simon and Shiraishi 1990; Minger et al. 1998), although testing in focal ischemia models is limited. Following adverse effects of NMDA antagonists in clinical trials, psychomimetic effects (e.g., hyperlocomotion, head weaving, and stereotyped sniffing) were studied in a small cohort of stroke rats and similar behavioral effects noted after administration (Loscher et al. 1998).

Gavestinel (GV150526), a NMDA receptor glycine site antagonist (Lees et al. 2000; Sacco et al. 2001) within 6 h of onset, showed no benefit in clinical outcome, mortality, or, in an imaging substudy, infarct attenuation in comparison with placebo (Warach et al. 2006). Gavestinel was tested in the Glaxo-Wellcome Research Laboratories, predominantly in permanent models of ischemia, and benefits on infarct measures of damage were found (Bordi et al. 1997; Di Fabio et al. 1999; Reggiani et al. 2001). Subsequent authors have hypothesized that NMDA receptor antagonists may hinder survival and regeneration (Hoyte et al. 2004); these outcomes were not typically measured in the studies.

#### AMPA Receptor Antagonists

Two clinical studies using YM872 (an AMPA receptor antagonist) within 6 h of stroke onset with and without concomitant rtPA, were prematurely terminated in 2003 after an interim futility analysis (Internet Stroke Center 2013). The study results have not been published. Animal studies have been undertaken in permanent and temporary occlusion models in rats (Kawasaki-Yatsugi et al. 1998; Shimizu-Sasamata et al. 1998; Haberg et al. 1998; Ni et al. 1998; Suzuki et al. 2003) and cats (Takahashi et al. 1998). While the overall result was positive, a number of individual trials showed minimal or no effect on infarct outcome in animal models.

#### 2.3.2 GABA Agonists

Studies using clomethiazole, a GABA(A) agonist (Wahlgren et al. 1999; Lyden et al. 2002), and diazepam, a GABA-ergic agent (Lodder et al. 2006) within 12 h of stroke onset showed no clinical benefit in comparison with placebo. In contrast, preclinical studies of clomethiazole in rats and marmoset models of stroke conducted by AstraZeneca and others tended to show a reduction in infarct volume and edema compared to control groups (Baldwin et al. 1994; Sydserff et al. 1995, 1996, 2000; Green and Cross 1994; Snape et al. 1993; Marshall et al. 1999, 2000). Studies of diazepam in animal stroke models are few: Diazepam did not reduce infarct volume in a filament occlusion model of stroke (Kuhmonen et al. 2002) but did so in a photothrombotic model (Aerden et al. 2004).

#### 2.3.3 Magnesium

The Intravenous Magnesium Efficacy in Stroke Trial (IMAGES) that randomized 2,589 patients within 12 h of stroke onset showed no clinical benefit for IV magnesium in comparison with placebo with a trend to higher mortality in the IV magnesium-treated group (Muir et al. 2004). The median time to treatment was 7 h, and only 3 % were treated within 3 h. The Field Administration of Stroke Therapy – Magnesium (FAST-MAG) Trial is currently recruiting to evaluate the safety and effectiveness of paramedic initiation of IV magnesium within 2 h of stroke onset (Internet Stroke Center 2013).

Magnesium has been relatively well characterized in investigator-sponsored trials in animal models of ischemia (Marinov et al. 1996; Lin et al. 2002; Westermaier et al. 2003; Meloni et al. 2006; O'Collins et al. 2011). It has been suggested that the effect of magnesium is via its effect on temperature, for where spontaneous hypothermia was prevented, no protective effect was seen (Zhu et al. 2004; Song et al. 2012; Campbell et al. 2008). A number of studies have found that magnesium is more effective when combined with other neuroprotective strategies (Schmid-Elsaesser et al. 1999; Zhao et al. 2010; Chung et al. 2004; Meloni et al. 2009; Campbell et al. 2008; Song et al. 2012; Wang et al. 2012), but not always (Lee et al. 1999). It has been argued that the sulfate salt is possibly more effective than the chloride salt (Izumi et al. 1991).

# 2.3.4 Antioxidants

# NXY-059

NXY-059, a nitrone with free radical scavenging properties, was subjected to more extensive preclinical testing than most other agents prior to entering clinical trials in humans although subsequently several deficiencies were identified (Savitz 2007). The SAINT I (Lees et al. 2006) and SAINT II (Shuaib et al. 2007) studies randomized 1,722 and 3,306 patients, respectively, to receive a 72-h infusion of NXY-059 commencing within 6 h of stroke onset. The SAINT I trial reported a small but statistically significant shift of the modified Rankin score (mRS) at 3 months, but there was no change in the Barthel index or NIHSS scale. The SAINT II trial failed to confirm the data reported in SAINT 1. When taken together, the investigators concluded that NXY-059 was ineffective for the treatment of acute ischemic stroke.

NXY-059 is interesting from a preclinical perspective in that it was one of the first drugs said to have been conducted according to rigorous experimental standards (the STAIR guidelines) (Kuroda et al. 1999; Marshall et al. 2001; Zhao et al. 2001; Lapchak et al. 2002a, b; Sydserff et al. 2002; Yoshimoto et al. 2002a, b; Lapchak and Araujo 2003; Marshall et al. 2003a, b; Lapchak et al. 2004). There was much hope that it would succeed at clinical trial, and after this failed, it received criticism for falling short in terms of methodological quality (Macleod et al. 2008; Bath et al. 2009; Savitz 2007). Nevertheless, the preclinical literature is still stronger than many other drugs which have been taken to clinical trial.

#### Tirilazad

Tirilazad, an aminosteroid that inhibits free radical-induced lipid peroxidation, showed no clinical benefit in acute stroke when initiated within 6 h of stroke onset (The RANTTAS Investigators 1996). A subsequent meta-analysis revealed that tirilazad increased death and disability by about one-fifth when given to patients with acute ischemic stroke (Tirilazad International Steering Committee 2000). In animal models, tirilazad (U74006F) has reduced infarct size in a rat and rabbit model of stroke but not cat (Alessandri et al. 2000; Beck and Bielenberg 1991; Hellstrom et al. 1994; Orozco et al. 1995; Oktem et al. 2000; Park and Hall 1994; Schmid-Elsaesser et al. 1998, 1999; Scholler et al. 2004; Takeshima et al. 1994; Umemura et al. 1994; Wilson et al. 1992; Xue et al. 1992; Zausinger et al. 2003a, b; He et al. 1993).

# 2.3.5 Membrane-Stabilizing Agents

#### Citicoline

Citicoline is an exogenous form of cytidine 5'-diphosphate choline, which is essential for the biosynthesis of membrane phospholipids. The ICTUS study (Dávalos et al. 2012) that randomized 2,298 patients to receive citicoline or placebo within 24 h of stroke onset did not confirm the clinical benefits that had been suggested by a meta-analysis of four smaller trials using this agent (Dávalos et al. 2012). In animal models of stroke, citicoline is one of the best agents characterized,

second only to tissue plasminogen activator (Schabitz et al. 1996; Onal et al. 1997; Andersen et al. 1999; Shuaib et al. 2000; Katsumata et al. 2001; Ataus et al. 2004; Hurtado et al. 2005, 2007; Adibhatla et al. 2006; Ramos-Cabrer et al. 2011; Xu et al. 2011; Diederich et al. 2012).

Piracetam, a GABA derivative, is also used clinically to improve cognitive function. Although taken to clinical trial, only a few studies have reported its function in animal models (Tortiglione et al. 2002; Muley et al. 2012). Repinotan (BAY x 3702) is a selective 5-HT1A receptor agonist developed and tested in animal models by Bayer Health Care (Berends et al. 2005) and tested in focal models (Semkova et al. 1998; Mauler and Horvath 2005).

#### 2.3.6 Calcium Modulators

#### Voltage-Gated Calcium Antagonists

A meta-analysis published in 2001 of 29 studies that included 7,665 patients with acute ischemic stroke treated with calcium antagonists reported no benefit with respect to clinical outcome or mortality (Horn and Limburg 2001). The calcium antagonists used were nimodipine in 23 studies, flunarizine in 3 studies, and isradipine and nicardipine in 1 study each. In 25 studies, including the Phase III TRUST (1,215 patients) and the American Nimodipine study (1,064 patients), treatment was started up to 48 h after stroke onset. There was no beneficial effect observed in the subgroup (660 nimodipine, 619 placebo) that received oral nimodipine within 12 h. Subsequent to this meta-analysis, the Very Early Nimodipine Use in Stroke (VENUS) trial reported the use of oral nimodipine within 6 h of stroke onset. This study was prematurely terminated by the steering committee because of sample size concerns. At the time 454 patients had been randomized and nimodipine was not associated with reduction of death or dependency at 3 months.

In animal studies, nimodipine was one of the earliest drugs trialed. More recently, it has been used a positive control against newer treatments, especially in studies of traditional Chinese medicines (Tian et al. 2005; Lu et al. 2011; He et al. 2012). It has been tested in mice, rats, and cats (Barnett et al. 1986; Gotoh et al. 1986; Hakim 1986; Sauter and Rudin 1986; Germano et al. 1987; Berger and Hakim 1988; Sauter et al. 1989; Jacewicz et al. 1990; Sakaki et al. 1991; Prehn et al. 1993; Herz et al. 1996; Cramer and Toorop 1998; Kawaguchi et al. 1999; Sobrado et al. 2003; Zausinger et al. 2003). In the 36 experimental comparisons included here, the median time for treatment was 5 minutes prior to ischemia.

#### Ginsenoside

Ginsenoside is a receptor-mediated calcium channel antagonist that does not lower blood pressure like voltage-gated calcium antagonists such as nimodipine discussed above. Liu et al. (2012) randomized 390 patients with acute ischemic stroke in a 3:1 ratio to receive a 14-day intravenous infusion of ginsenoside or placebo within 72 h after the onset of stroke. The primary outcome was positive with improved distribution of disability scores on the modified Rankin scale (mRS) at 90 days in the group receiving ginsenoside. This is a promising result that needs verification in a larger RCT, bearing in mind that the initial NXY-059 trial reported a similar positive result based on shift of the mRS scores that was not confirmed in a second larger trial.

Ginsenosides (ginseng saponins) are the active component isolated from the traditional Chinese medicine ginseng, from the root of Panax ginseng. Over 30 different ginsenosides have been isolated, a number of these have been recently studied in animal models: ginsenoside Rb(1), ginsenoside Rd, ginsenoside Rh2, ginsenoside Rg(2), ginsenoside RG3, and fermented ginseng (Bae et al. 2004). Animal models suggest that the mode of action is on mitochondrial energy metabolism and altered activity of the mitochondrial permeability transition pore (Tian et al. 2009) and potentially via its effect on neural plasticity (Cheng et al. 2005) and angiogenesis (Yang et al. 2012). Rodent models have shown dose-dependent reductions in infarct size; however, sample sizes tend to have been small, and in many studies animals were treated prior to ischemia (Park et al. 2004; Tian et al. 2005; Zhang et al. 2006; Lu et al. 2011; Ye et al. 2011a, b, c; He et al. 2012; Yang et al. 2012; Zhu et al. 2012).

# 2.3.7 Anti-apoptotic and Trophic Factors

#### Cerebrolysin

First developed in Russia in the 1970s, cerebrolysin is a porcine brain-derived preparation of low-molecular-weight neuropeptides (10 kDa) and free amino acids that shows pharmacodynamic properties similar to those of naturally occurring neurotrophic factors. The CASTA study investigators randomized 1,070 patients to receive cerebrolysin or placebo within 12 h of stroke onset (Heiss et al. 2012). There was no difference in the primary outcome of a global functional scale, but there was a trend towards lower disability and reduced mortality in the cerebrolysin group for patients with more severe stroke (NIHSS >12). Cerebrolysin has demonstrated efficacy in a limited set of animal focal ischemia experiments (Schwab et al. 1998; Ren et al. 2007; Hanson et al. 2009; Zhang et al. 2010).

Erythropoietin, either alone or in combination with another agent after ischemic stroke, has attracted much interest in the last decade. Where animal tests have been undertaken, the time window for delivery tends to have been longer than other potential neuroprotectives and more emphasis has been placed on behavioral tests (Bernaudin et al. 1999; Brines et al. 2000; Wang et al. 2004, 2007; Esneault et al. 2008; Belayev et al. 2009; Fletcher et al. 2009; Jia et al. 2010; Zechariah et al. 2010).

#### 2.4 Possible Reasons for Failure of Neuroprotection

With such positive data from animal experiments, how is it then that the promise of neuroprotection has thus far failed to translate into clinical benefit in human stroke studies (Ginsberg 2008; Sutherland et al. 2012)? There are numerous

preclinical and clinical reasons that could account for this failure. Failure of the initial clinical trials led to the development of the STAIR guidelines in 1999 for preclinical animal testing (STAIR Group 1999). In the wake of negative human clinical trials of compounds such as NXY-059 that had appeared promising in animal trials that fulfilled these guidelines, the criteria were revised in 2009 (Fisher et al. 2009).

A key issue in the human clinical trials is whether the agent actually reached the target tissue, the ischemic penumbra. The time window for effective thrombolysis in humans is 4.5 h after sstroke onset with greatest efficacy at earlier time points when the penumbra is maximal. By contrast, the completed Phase II/III studies of neuroprotective agents in humans had time windows of up to 48 h although efficacy was demonstrated at a shorter time window in the corresponding animal studies where there is no evidence of effective neuroprotection beyond 6 h. In the animal data surveyed above, the median time of treatment administration was 10 min after stroke onset.

Ischemic stroke in humans is heterogeneous with a multitude of subtypes and mechanisms. In contrast, animal studies have used relatively homogenous MCA occlusion models, but this particular stroke subtype accounts for only about 25 % of ischemic strokes in humans. Potential therapies have been tested almost exclusively in male animals (unless a female hormone is implicated in the mode of action) and in young animals (our data suggests that less than 1 % of testing occurs in middle-aged or old animals).

Neurotoxicity or neuroprotection may also occur as a covariate in animal models, e.g., anesthetic agents such as ketamine have effects on regional cerebral blood flow, EEG, sensation, and memory. Finally, there are important anatomical differences in gray and white matter distribution animal and human brains. The ischemic cascade has been studied primarily in gray matter that accounts for 90 % of rodent brain, but white matter accounts for approximately 50 % of human brain tissue involved by ischemic stroke (Ho et al. 2005).

# 3 The Future of Neuroprotection in Ischemic Stroke

#### 3.1 Current Trends

The large numbers of negative clinical trials of neuroprotection in humans have led to calls for abandoning future trials, citing ineffectiveness of this strategy (The Lancet 2006). Others have reasonably argued for improved design of future trials of neuroprotectants in humans, primarily with shorter time windows and use in conjunction with thrombolytic therapy (Ginsberg 2008; Sutherland et al. 2012). Current ongoing trials of neuroprotection strategies in ischemic stroke incorporating these changes include intravenous magnesium within 2 h of onset (Internet Stroke Center 2013) and uric acid, an endogenous antioxidant in conjunction with rtPA (Amaro et al. 2010). In preclinical development, current trends in neuroprotection include a focus on the microvasculature, inflammation, and the neurovascular unit and, secondly, a focus on endogenous neuroprotection and regeneration (stem cells, hormones, oxygen metabolism).

### 3.2 Improved Drug Development Procedures

Few of the agents tested in animal models of stroke have been developed specifically for treatment of ischemic stroke. This will reflect a bias against publication of potentially patentable drugs. However, it also suggests a need for better funding of stroke target identification and validation, together with better development of lead drug candidates before taking candidates to preclinical studies.

# 3.3 Toxicity in Animal Models of Stroke

Although toxicity is a frequent cause of drug failure at clinical trial, preclinical investigation of toxicity is generally looked at further down the preclinical pathway. Simultaneous optimization of drug efficacy and ADMET properties (absorption, distribution, metabolism, elimination, toxicology) is a strategy which could be looked at in greater detail in stroke drug development. To this end, improved computational tools for better prediction of toxicity based on biochemical pathways and drug characteristics may assist.

Expanding preclinical outcome measures beyond the standard measures of brain damage and neurological score may help identify potential neurotoxicity of drugs earlier. Cognitive deficit which has been noted following administration of NMDA antagonists could have been detected in animal models with tests typically deployed in other areas of neuroscience, for instance, locomotion (hyperlocomotion), prepulse inhibition disturbances, and disruptions to social interactions. Side effects of stroke drugs can include neurotoxicity but may also extend to genotoxicity, carcinogenicity, reproductive and developmental toxicity, and skin sensitization or irritation.

# 4 Conclusion

Ischemic stroke is a complex disease that straddles the vascular and nervous systems and is frequently confounded by comorbidities such as hypertension, diabetes and age-related degeneration. Thus, identifying and targeting individual causes of neurotoxicity in stroke is a great challenge for the clinical testing and laboratory development of novel therapeutics. The clinical and animal trials reviewed here have tended to focus on protecting the brain against damage caused by free radical scavengers and excitoxicity, or have used pharmacological means to target calcium modulation, membrane stabilization and trophic factors.

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# **Neurotoxicity in Huntington Disease**

# C. T. Loy and A. J. Hannan

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C.T. Loy (🖂)

School of Public Health, The University of Sydney, Sydney, Australia

A.J. Hannan The Florey Institute of Neuroscience and Mental Health, The University of Melbourne, Melbourne, Australia e-mail: anthony.hannan@florey.edu.au

Huntington Disease Service, Westmead Hospital, Sydney, Australia e-mail: clement.loy@sydney.edu.au

#### Abstract

Huntington Disease (HD) is an autosomal dominant, progressive, neurodegenerative disorder, due to abnormal CAG expansion in the Chromosome 4 *Huntingtin (HTT)* gene. Typical age of onset is in the 30s-40s, and symptoms include involuntary movements, cognitive/behavioural symptoms and psychiatric disorders. As a monogenic disorder, the aetiology of HD is well established however our understanding of the pathogenesis of HD is still evolving. In particular, mutation in the *HTT* gene confers neurotoxicity on multiple levels, ranging from the genetic/protein level, intracellular level, intercellular level, to the end-organ level. On a genetic/protein level, a *HTT* mutation is associated with toxic gain of function, loss of normal function and altered transcription of other genes. On an intracellular level, it is associated with mitochondrial dysfunction and impaired cytoskeleton/intra-cellular trafficking. On an intercellular level, it is associated with impaired synaptic transmission, excitoxicity and inflammation. On an organ level, both the brain and peripheral tissues are affected. We will review each of these areas in this article.

#### **Keywords**

Huntington disease • Molecular genetics • Neurodegeneration • Neurotoxicity • Polyglutamine • Tandem repeat disorder

## 1 Introduction

Huntington Disease (HD) is an autosomal-dominant, progressive, neurodegenerative disorder, due to abnormal CAG expansion in the chromosome 4 *Huntingtin (HTT)* gene. It is one of the most common neurogenetic disorders (MacMillan and Harper 1991), and has devastating consequences for patients and their families (Vamos et al. 2007). Typical age of onset is in the 30s–40s, and symptoms include involuntary movements, cognitive/behavioral symptoms, and psychiatric disorders (Walker 2007). While symptomatic treatments exist, these do not prolong survival, and median survival is about 20 years (Walker 2007). HD strikes at a productive period in work and family life, and requires a high level of complex care. Consequently, the direct and indirect cost of HD has been informally estimated to be two billion dollars per year, in the United States alone (Society for Neuroscience 2005).

## 2 Overview

As a monogenic disorder, the etiology of HD is well established (The Huntington's Disease Collaborative Research Group 1993), however our understanding of the pathogenesis of HD is still evolving. In particular, mutation in the *HTT* gene appears to confer neurotoxicity on multiple levels, ranging from the genetic/protein level, intracellular level, intercellular level, to the end-organ level (Table 1). We will review each of these areas in turn.

<b>Table 1</b> Mutation in the Huntingtin gene leads to neurotoxicity on multiple levels		Consequences of mutation in the <i>Huntingtin</i> gene
	Neurotoxicity on a genetic/ protein level	Toxic gain of function:
		1. Altered conformation
		2. Aberrant posttranslational modifications
		3. Impaired clearance
		4. Toxic proteolytic cleavage products
		5. Increased propensity to aggregate
		Loss of normal function
		Altered transcription of other genes
	Neurotoxicity on an intracellular level	Role of the intranuclear inclusion body
		Mitochondrial dysfunction
		Impaired cytoskeleton and intracellular trafficking
	Neurotoxicity on an intercellular level	Impaired synaptic transmission
		Excitotoxicity and the kynurenine pathway
		Inflammation (?)
	Neurotoxicity on the end- organ level	Brain
		Peripheral tissues

## 3 Neurotoxicity on a Genetic/Protein Level

The *HTT* gene is located on the short arm of chromosome 4 (4p16.3) (Meyer et al. 2012). It has 67 exons and encodes a 348-kDa nuclear protein which is widely expressed in the human brain as well as other tissues (Strong et al. 1993). Exon 1 of the *HTT* gene contains a polymorphic CAG repeat, which is expanded among people with HD. The general population carries up to 26 repeats at this locus, and people carrying 40 or more repeats almost invariably develop clinical HD (ACMG/ASHG 1998). People with 36–39 CAG repeats have reduced penetrance. People with 27–35 CAG repeats do not develop clinical HD, but their offspring may do so, due to meiotic instability of the CAG repeat and further expansion across generations (Leeflang et al. 1995).

## 3.1 Toxic Gain of Function

From a genetic point of view, the prevailing hypothesis is that the expanded CAG repeats lead to a toxic gain of function. This is supported by: (1) the dominant mode of inheritance, (2) association of longer CAG repeats with earlier age of onset (Lee et al. 2012), and (3) no worsening of clinical HD phenotype among people

without a normal copy of *HTT* gene (i.e., people with homozygous HD mutations) (Wexler et al. 1987). In addition, a transgene expressing the human *HTT* Exon 1 with expanded CAG repeats was sufficient to cause a HD-like phenotype in the R6 lines of transgenic mice (Mangiarini et al. 1996).

Accordingly, the mutant huntingtin protein (mHTT) has been studied to elucidate a potential pathogenic mechanism for HD. Compared to wildtype huntingtin protein (wHTT), mHTT has (1) altered conformation, (2) aberrant posttranslational modifications, (3) impaired clearance, (4) toxic proteolytic cleavage products, and (5) increased propensity to aggregate (Ross and Tabrizi 2011).

## 3.1.1 Altered Conformation

In wHTT, the CAG expansion translates to an N-terminal polyglutamine tail which is flexible and can adopt a variety of conformations (Kim et al. 2009). However, when the CAG region is expanded into the pathogenic range, mHTT develops a compact two-strand hairpin conformation (Peters-Libeu et al. 2012). This hairpin epitope was identified using the 3B5H10 antibody, which also predicts the severity of neurodegeneration upon live cell imaging (Miller et al. 2011). Exactly how this hairpin structure leads to neurodegeneration is currently unclear, although early pathogenic processes are likely to include changes in protein solubility, aggregation, and protein–protein interactions.

#### 3.1.2 Aberrant Posttranslational Modifications

Posttranslational modification (PTM) of mHTT is altered compared to wHTT. Examples of these PTM include phosphorylation, palmitoylation, ubiquitination, and acetylation. Huntingtin is constitutively phosphorylated at multiple sites (Schilling et al. 2006), and in general, mHTT is less phosphorylated than wHHT (Ehrnhoefer et al. 2011). For instance, in cell models, phosphorylation of Threonine-3 in HTT declines in a dose-response manner, with increasing CAG repeat length (Aiken et al. 2009). When phosphorylation was mimicked using sitespecific mutagenesis for Serines-13 and -16, mHTT aggregation was reduced in vitro, and motor/psychiatric symptoms improved in a mouse model (Gu et al. 2009). Palmitoylation at Cysteine-214 of the HTT protein is also reduced, in the presence of a pathogenic CAG repeat (Yanai et al. 2006). Cells expressing mHTT resistant to palmitoylation are associated with faster inclusion body formation (Yanai et al. 2006). Ubiquitination is associated with protein targeting for proteasome degradation. Mutant HTT with expanded glutamine repeats have a slower rate of degradation in the proteasome (Jana et al. 2001). In addition, a protein specific to the striatum, the Ras Homologue Enriched in the Striatum (Rhes) protein, selectively increases SUMO-ylation (Small Ubiquitin-like modifier) of mHTT. This results in disaggregation of mHTT and cytotoxicity, and is consistent with the anatomical distribution of HTT (Subramaniam et al. 2009). Finally, acetylation at Lysine-444 of mHTT is associated with tagging for autophagy in vitro. This reverses toxic effects of mHTT in C. elegans and cell models, and is an example where alteration in posttranslational modification of mHTT is beneficial rather than deleterious (Jeong et al. 2009).

#### 3.1.3 Impaired Clearance

Clearance of mHTT is impaired. In addition to the slower rate of degrading mHTT in the proteasome, N-terminal fragments of mHTT were found to directly impair functioning of the Uniquitin/Proteosome System (UPS) in a cellular model (Bence et al. 2001). More recently, this has been confirmed in vivo, using a UPS reporter/inducible HD mouse model (Ortega et al. 2010). UPS dysfunction was found transiently after expression of N-terminal mutant HTT, and recovered with appearance of inclusion bodies.

## 3.1.4 Toxic Proteolytic Cleavage Products

A number of experiments support the notion that neurotoxicity in HD arises from toxic proteolytic cleavage products of mHTT rather than the full mHTT molecule. These include: (1) sufficiency of *HTT* Exon 1 to cause an HD-like phenotype in a transgenic mouse model (Mangiarini et al. 1996), and (2) normal phenotype in a mouse model expressing caspase-6 resistant mHTT (Graham et al. 2006). In addition to caspases, calpains also play a role in proteolytic cleavage (Southwell et al. 2011). More recently, CAG repeat length-dependent aberrant splicing of the *HTT* gene has been found to be an alternative mechanism in generating N-terminal mHTT fragments (Sathasivam et al. 2013).

## 3.1.5 Increased Propensity to Aggregate

Early neuropathologic studies of the brains of people with HD found intranuclear inclusion bodies that contained protein aggregates immunoreactive to antibodies raised to the N-terminal 17 amino acids, but not an internal epitope for HTT (Becher et al. 1998). This raised the question as to whether these N-terminal HTT fragments, aggregated in inclusion bodies, were pathogenic. As evidence was generated showing that the inclusion bodies could be protective rather than pathogenic (Arrasate et al. 2004) (see Sect. 4.1 for further details), attention was shifted to polyglutamine monomers and oligomers, which may be more neurotoxic (Miller et al. 2011). Oligomer formation is faster and more extensive for HTT fragments with longer polyglutamine tracts (Legleiter et al. 2010). This increased propensity for mHTT to aggregate also provides an explanation for the inverse relationship between CAG repeat length and age of onset in HD. Interestingly, an aggregation kinetic study has found a static population of oligomers despite ongoing conversion of mHTT monomers to inclusion bodies in a cell models (Olshina et al. 2010). This suggests that neurotoxicity due to oligomers may be a constant process, and correlates poorly with inclusion body formation.

## 3.2 Loss of Normal Function

While the prevailing hypothesis for neurotoxicity of CAG expansion is a toxic gain of function, there is also an argument that loss of normal function may play a role (Cattaneo et al. 2005). Mouse models without a normal copy of the murine

*HTT* analogue gene are embryonically lethal (Duyao et al. 1995; Nasir et al. 1995). Inactivation of the murine *HTT* analogue genes postnatally also leads to a progressive neurodegenerative phenotype (Dragatsis et al. 2000). These underline the role of wHTT in normal neuronal functioning, which include: (1) upregulation of brain-derived neurotrophic factor (BDNF) (Zuccato et al. 2001), (2) interaction with the motor proteins and microtubules in BDNF transport (Gauthier et al. 2004), and (3) interaction with literally hundreds more proteins (Kaltenbach et al. 2007), including ones important for synaptic function (Milnerwood and Raymond 2010).

#### 3.3 Altered Transcription of Other Genes

Wildtype HTT upregulates BDNF by sequestering repressor element-1 transcription factor/neuron restrictive silencer factor (REST/NRSF) in the cytoplasm, reducing its availability to bind neuron restrictive silencer element (NRSE) in the nucleus, thus allowing increased transcription (Zuccato et al. 2003). This mechanism also applies to a number of other NRSE-containing genes, and is impaired in mHTT. NRSE is only one of many pathways through which gene transcription is altered in HD. Other examples include nuclear receptor compressor (N-CoR) (Boutell et al. 1999), transcription factor Sp1 (Dunah et al. 2002), the Histone Acetyltransferase cAMP response element-binding protein (CREB) (Nucifora et al. 2001), and RNA-mediated gene silencing through association with the Argonaute (Savas et al. 2008). As a consequence, mHTT is associated with widespread alteration in gene transcription both in development (Feyeux et al. 2012) and diseased brains (Hodges et al. 2006). Drugs targeting some of these transcriptional changes, e.g., Histone Deacetylase Inhibitors, appear to be promising therapeutic agents (Steffan et al. 2001).

#### 4 Neurotoxicity on an Intracellular Level

## 4.1 Role of the Intranuclear Inclusion Body

A characteristic feature of the neuropathology HD is the intranuclear inclusion body, which stains positive for ubiquitin and HTT. They also stain positive for antibodies raised to the N-terminal 17 amino acids, but not those raised to an internal epitope of HTT. This was the case in both the R6/2 mouse model (Mangiarini et al. 1996) and human archival tissue (Becher et al. 1998). The role of inclusion bodies in pathogenesis was clarified with automated microscopy and live cell imaging, where inclusion body formation was associated with improved neuron survival rather than death (Arrasate et al. 2004). Since then, the focus on mHTT has shifted to mHTT monomers and oligomers.

#### 4.2 Mitochondrial Dysfunction

Mutant HTT can impair cellular mitochondrial function directly and indirectly. Direct effects of mHTT on the mitochondria include altered calcium handling and mitochondrial dynamics. Mitochondrial capacity to retain calcium is impaired in HD (Panov et al. 2002), where mHTT is thought to interrupt membrane integrity (Choo et al. 2004; Rockabrand et al. 2007). This, in turn, adds to the effect of glutamate/calcium-mediated excitotoxicity. Mutant HTT contributes toward increased intracellular calcium by sensitizing activation of the type 1 inositol-(1,4,5)-triphosphate receptor, thus increasing calcium release from the endoplasmic reticulum (Tang et al. 2003).

Mitochondria undergo continual fissure and fusion, and alteration of its dynamics can lead to neurodegeneration (Knott et al. 2008). Mutant HTT triggers mitochondrial fragmentation and arrests mitochondrial transport, through abnormal interaction with the GTPase dynamin-related protein 1 (DRP1) (Song et al. 2011). Interestingly these cellular changes can be rescued by reducing DRP1 activity.

Finally, mHTT also impairs mitochondrial function indirectly by inhibiting expression of peroxisome proliferator-activated receptor [PPAR]-gamma coactivator-1 alpha (PGC-1 alpha), a transcriptional coactivator that regulates mitochondrial function (Cui et al. 2006). The mechanism for this is likely to be through interference with the CREB transcription pathway.

## 4.3 Impaired Cytoskeleton and Intracellular Trafficking

Wildtype HTT binds to monomeric tubulin and microtubules, while mHTT disrupts integrity of microtubule filaments (Trushina et al. 2003). Mutant HTT is associated with neurite retraction in a cell culture model, indicating cytoskeletal collapse. Toxicity of mHTT can also be modulated by microtubule stabilization using low-dose taxol (Trushina et al. 2003).

Mutant HTT is associated with impaired axonal transport, with possible mechanisms including: titration of motor proteins such as kinesin from other cargoes (Gunawardena et al. 2003), direct blockage of axonal transport by aggregates (Lee et al. 2004), and by reducing kinesin-1 binding to microtubules through mHTT associated cJun N-terminal kinase (JNK) activation (Morfini et al. 2009).

Loss of normal function of wHTT, as an integrator of molecular motors, may also contribute toward the disrupted intracellular trafficking in HD (Caviston and Holzbaur 2009). An example of one such molecular binding partner for HTT is Huntingtin-Associated Protein 1 (HAP1) (Li et al. 1995). HAP1 is concentrated in axon terminals and promotes neurite extension (Li et al. 2000). Mutant HTT reduces HAP1 binding to molecular motors, resulting in decreased level of the nerve growth factor receptor Tropomyosin-receptor kinase A (TrkA), and neurite outgrowth (Rong et al. 2006). Translating to human studies, an amino-acid substituting polymorphism in *HAP1* was found to be associated with a later age of onset in HD (Metzger et al. 2008). Finally, mHTT also impairs transport of mitochondria (Song et al. 2011) and vesicles (Gauthier et al. 2004).

## 5 Neurotoxicity on an Intercellular Level

#### 5.1 Impaired Synaptic Transmission

In addition to vesicular transport, synaptic transmission is also affected in HD (Smith et al. 2005). In exocytosis, mHTT reduces the level of Complexin II, a protein involved in the transport vesicle/presynaptic protein-binding process (Edwardson et al. 2003). In that cell model, there was a corresponding reduction in calcium-triggered exocytosis of neurotransmitters.

Mutant HTT impairs endocytosis through interaction with caveolin-1, which has a role in forming invaginations of the plasma membrane (Trushina et al. 2006). There is some suggestion that mHTT could impair clathrin-mediated endocytosis; however, the data is conflicting (Trushina et al. 2006; Borgonovo et al. 2013).

These, together with other factors, have resulted in complex changes of glutamate and dopamine levels in the brain (Andre et al. 2010). Another neuro-transmitter affected by mHTT is Gamma-AminoButyric Acid (GABA) (Pearson and Reynolds 1994).

### 5.2 Excitotoxicity and the Kynurenine Pathway

The association between excitotoxicity and HD was made prior to molecular characterization of HTT. This was based on the shared, extrinsic axon sparing phenotype, between the human HD brain and lesion models using excitatory amino acids (Schwarcz et al. 2010). In particular, the Quinolinic acid (QUIN) lesioned rats were found to have similar neurochemical profiles to human HD brains. These neurochemical profiles were characterized by depletion of GABA and Substance P, and relative sparing of Somatostatin/Neuropeptide Y neurons (Beal et al. 1986). QUIN level was also raised in early stage HD brains (Guidetti et al. 2004). QUIN is a downstream product of the kynurenine pathway (KP), which catabolizes the amino-acid tryptophan. It forms part of the neuroinflammatory response and acts as an excitotoxin through the *N*-methyl-D-aspartate (NMDA) receptor (Guillemin 2012). Recent interest in the KP for HD has grown, following reports that modulation of the KP led to improved life span in a mouse model (Zwilling et al. 2011), and a *Drosophila* model of HD (Campesan et al. 2011).

## 5.3 Inflammatory Response

There are a few lines of evidence, in addition to QUIN, that suggest a role of inflammation in HD (Soulet and Cicchetti 2011). In a mouse model where mHTT is selectively expressed in astrocytes only, the phenotype remained similar to the corresponding fully expressed HD mouse, albeit with a late onset (Bradford et al. 2009). Microglial activation measured on Positron Emission Tomography scans also correlated with clinical severity of people with HD (Pavese et al. 2006). Nonetheless, the exact role of inflammation in the pathogenesis of HD remains unclear.

## 6 Neurotoxicity on an Organ Level

## 6.1 Brain

Since Graveland et al.'s Golgi impregnation study in 1985, medium-sized spiny neurons in the striatum have been considered the primary cell type affected in HD (Graveland et al. 1985). Medium-sized spiny neurons require a high level of energy to maintain their unusually high transmembrane potentials. This makes them more vulnerable than other cell-types to the disrupted cellular respiration in HD (Mitchell et al. 1999).

However, the striatum is not the only brain region affected in HD. Recently, a large HD brain autopsy series (N > 500) (Hadzi et al. 2012) highlighted the role of the cerebral cortex. Forty-one neuropathologic ratings from various regions of the brain were categorised into two clusters: the striatal and cortical clusters. Neurodegeneration in the striatal cluster had a stronger correlation with HD CAG repeat size, but the cortical cluster had a stronger correlation with brain weight. These suggested that neurodegeneration in the striatum and the cortex do not necessarily proceed in a coupled manner.

#### 6.2 Peripheral Tissues

Finally, the effect of mHTT reaches beyond the brain (Sassone et al. 2009; van der Burg et al. 2009). For instance, derangement in the hypothalamic–pituitary–adrenal axes is not the only potential mechanism for weight loss in HD. Other contributory mechanisms include the effect of mHTT on the gastrointestinal tract (impaired motility and malabsorption) (van der Burg et al. 2011), pancreas (altered glucose homeostasis) (Lalic et al. 2008), adipose tissue (Phan et al. 2009), and a hypermetabolic state (Mochel et al. 2007).

## 7 Conclusion

Given the complex and varied mechanisms leading to neurotoxicity in HD, it is unlikely that a therapeutic agent based on any single mechanism will be able to reverse all disease manifestations of HD. This makes oligonucleotides (RNA interference and antisense oligonucleotides) (Sah and Aronin 2011) a particularly attractive therapeutic approach, as they will be acting upstream of most other neurotoxic mechanisms. Animal studies using oligonucleotides have been promising (Kordasiewicz et al. 2012; Yu et al. 2012), and the first round of human safety trials is keenly awaited. However, complementary therapeutic approaches, addressing a range of pathogenic targets, need to be addressed in parallel, to combat this devastating disease.

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# Neurotoxicity: A Complex Multistage Process Involving Different Mechanisms

Ewa Niedzielska, Bartłomiej Rospond, Lucyna Pomierny-Chamioło, Anna Sadakierska-Chudy, and Małgorzata Filip

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#### Abstract

Neurotoxicity is defined as alterations causing physical damage or disruption in functions of the cells (neurons, glial cells) of the nervous system. There are several mechanisms by which neurotoxic processes take place. These include high concentrations of glutamate and/or calcium ions, increased oxidative damage, and reactive species (RS) such as free radicals, mechanisms linked to execution of

Laboratory of Drug Addiction Pharmacology, Institute of Pharmacology Polish Academy of Sciences, Kraków, Poland

e-mail: malgorzata.filip@uj.edu.pl; filip@if-pan.krakow.pl

E. Niedzielska • B. Rospond • L. Pomierny-Chamioło

Department of Toxicology, Faculty of Pharmacy, Jagiellonian University, Kraków, Poland

A. Sadakierska-Chudy Laboratory of Drug Addiction Pharmacology, Institute of Pharmacology Polish Academy of Sciences, Kraków, Poland

M. Filip (🖂) Department of Toxicology, Faculty of Pharmacy, Jagiellonian University, Kraków, Poland

apoptosis, glial dysfunction injury, and/or genetic-epigenetic defects in protein clearance. Each of these mechanisms reciprocally modulates their own actions and is capable of neuronal cell death followed by nervous system dysfunction.

## Keywords

Apoptosis • Calcium ions • DNA damage • Epigenetic • Glia activation • Glutamate • Neurotoxicity mechanism • Neurotoxicity pathway • Oxidative stress

List of Abbreviations				
5-HT	Serotonin			
AIF	Proapoptotic inducting factor			
AMPA	α-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid			
Apaf-1	Apoptotic protease activating factor-1			
ATP	Adenosine triphosphate			
CAT	Catalase			
CNS	Central nervous system			
CpG	Cytosine-guanine dinucleotide			
CREB	cAMP response element binding			
CYP	Cytochrome P450			
DISC	Death-inducing signaling complex			
DNMT	DNA methyltransferase			
EAAT	Excitatory amino-acid transporter			
endo-G	Endonuclease G			
esiRNA	Endogenous small interfering RNA (endo-siRNA)			
FADD	Fas-associated death domain			
FasL	Fas ligand			
GABA	γ-aminobutyric acid			
GPx	Glutathione peroxidase			
GR	Glutathione reductase			
GSH	Glutathione			
GSSG	Glutathione disulfide			
HAT	Histone acetyltransferase			
HDAC	Histone deacetylase			
HDM	Histone demethylase			
HMT	Histone methyltransferase			
IAP	Inhibitor of apoptosis protein			
IL	Interleukin			
JNK	c-Jun NH <sub>2</sub> -terminal kinase			
lncRNA	Long noncoding RNA			
MAO	Monoamine oxidase			
MDA	Malondialdehyde			
miRNA	microRNA			
MMP	Metalloproteinase			
ncRNA	Noncoding RNA			
NMDA	<i>N</i> -methyl-D-aspartic acid			

NOS	Nitric oxide synthase
NSL	Nuclear localization signal
PARP	Poly-ADP ribose polymerase
piRNA	Piwi-interacting RNA
RBS	Reactive bromine species
RCS	Reactive chloride species
RIP	Receptor-interacting protein
RNS	Reactive nitrogen species
ROS	Reactive oxygen species
RS	Reactive species
SMAC	Small mitochondria-derived activator of caspase
SOD	Superoxide dismutase
SODD	Silencer of death domain
TNF	Tumor necrosis factor
TRADD	TNF receptor-associated death domain
TRAF2	TNF-R-associated factor 2
TRPM	Transient receptor potential melastin
VGLUT	Vesicular glutamate transporter
xCT	Cystine-glutamate antiporter

## 1 Introduction

Neurotoxicity is defined as alterations in the normal activity of the nervous system to cause its physical damage (e.g., cell death) or disruption in functions (e.g., neurocognitive deficits, poor judgments, and diminished decision-making capacity). Within the central nervous system (CNS), neuronal cells are key factors for neurotoxicity and degeneration in many diseases. However, in the mid-1990s research has shown that glial cells (i.e., astroglia, microglia, and oligodendrocytes) also play a number of active roles in the brain and are sensitive indicators for pathology of the CNS (see also Clark et al. 2013). Thus, astrocytes – apart from provision of nutrients to the nervous tissue, maintenance of both extracellular ion balance and the blood-brain barrier, and a role in the repair and scarring process of the brain and spinal cord following traumatic injuries – are engaged in the release or absorption of neural transmitters (called gliotransmission) (Ben Achour and Pascual 2012; Santello et al. 2012). For example, astrocytes propagate intercellular calcium waves over long distances in response to stimulation and, similar to neurons, release transmitters in a Ca<sup>2+</sup>-dependent manner (Zorec et al. 2012), while a new report indicates also that astrocytes from adult human brain are y-aminobutyric (GABA)-ergic cells possessing enzymes and receptors and modulating microglial activity representing GABAceptive cells (Lee et al. 2011). The concept of a "tripartite synapse" referring to the relationship occurring at synapses among a presynaptic and a postsynaptic element as well as a glial element has been introduced (Santello et al. 2012). On the other hand, microglial cells are activated due to neuronal damage leading to reactive gliosis and to cytoarchitectonic changes accompanied by alterations in surface receptor and channel expression as well as to release of neuroactive soluble factors (see below) that in turn exert actions on neuronal function and cell death. Recently, it was found that oligodendroglia in the CNS is not only linked to myelin formation and maintenance, but they actively participate in neuronal survival and development as well as in neurotransmission and synaptic function including neurotoxic processes (Deng and Poretz 2003).

Neurotoxicity can result from exposure to several factors (intrinsic and exogenous xenobiotics) that disturb the normal physiology and function of the nervous system. The exogenous and intrinsic factors influence each other's actions.

## 2 Intrinsic Factors Engaged in Neurotoxicity

There are several endogenous brain toxins which in high amount may lead to neurotoxicity. When present in high concentrations, glutamate, calcium ions, and reactive species (RS) such as free radicals, proinflammatory cytokines, tumor necrosis factor alpha (TNF- $\alpha$ ), or neurotrophins released by microglia are potentially neurotoxic (Aloisi 2001; Aloisi et al. 2001; Hanisch 2002). All of them reciprocally modulate their own actions and are responsible for neuronal cell death followed by nervous system dysfunction (e.g., loss of motor control; cognitive deterioration; loss of memory, vision, and/or intellect; uncontrollable obsessive and/ or compulsive behaviors; delusions; headache; or neurodegenerative diseases). The most common to damage cells within the CNS are glutamate, increased cell calcium concentrations, RS and increased oxidative damage, mitochondrial dysfunction linked to apoptotic pathways, glial dysfunction injury, and/or genetic-epigenetic defects in protein clearance (cf. Halliwell 2006).

#### 2.1 Neurotoxicity Induced by Glutamate

Glutamate serves a principal brain neurotransmitter, while its neurotransmission impairment such as high concentrations (> $\mu$ M in extracellular fluids) evokes excitotoxicity to neurons. The glutamate concentration equilibrium is extremely delicate and is under control of glutamate transporters that pump the glutamate back from the neuronal synapse into the glia and neurons and allow glutamate to be recycled for repeated release via glutamate-glutamine neuronal glial cycle.

There are two general classes of glutamate transporters, those that are dependent on a sodium ion gradient (so-called the excitatory amino-acid transporters; EAATs) and those that are not dependent (vesicular glutamate transporters (VGLUTs) and cystineglutamate antiporter (xCT)). EAATs are also dependent on transmembrane K<sup>+</sup> and H<sup>+</sup> concentration gradients and are known also as "sodium- and potassium-coupled glutamate transporters" or "high-affinity glutamate transporters." In humans and rodents, five subtypes of EAATs have been identified: subtypes EAAT1–2 are found in membranes of glial cells (astrocytes, microglia, and oligodendrocytes) and in endothelial cells, EAAT3–4 are located on neurons, while EAAT5 is found in the retina. Four VGLUTs (VGLUTs 1–3 and sialin) are found in the membrane of glutamate-containing synaptic vesicles. The xCT located to the plasma membrane of cells prefers cystine and glutamate as its substrates in the way that the anionic form of cystine is transported in exchange for glutamate to glial cells. The xCT protein is upregulated in glial cells upon oxidative stress and plays an essential role to protect neurons. The glial transporters – in particular the various splice variants of EAAT2 and xCT – play the largest role (>90 %) in regulating extracellular glutamate levels. Data suggest that astrocytes also signal to neurons through Ca<sup>2+</sup>-dependent release of glutamate (Hua et al. 2004).

In the event of any disruption in the function of glutamate transporters, a high concentration of glutamate accumulates at the glutamate receptors. Exposing neurons to high glutamate concentrations results in the activation of ionotropic glutamate receptors such as the *N*-methyl-D-aspartic acid (NMDA) receptors or  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) receptors, which induce faster mitochondrial Ca<sup>2+</sup> uptake (Rego and Oliveira 2003). NMDA receptor overstimulation allows high levels of calcium ions to enter to cells, causing formation of mitochondrial RS and lipid peroxidation (Maciel et al. 2001). Moreover, extrasynaptic NMDA receptor activation promotes a dominant CREB (cAMP response element binding) dephosphorylation pathway that leads to protective CREB function shutoff causing loss of mitochondrial membrane potential and finally apoptosis (Hardingham and Bading 2010).

It should be added that NMDA glutamatergic receptors are stimulated also by quinine, being a product of the autoxidative pathway of dopamine metabolism (Lieb et al. 1995) which was shown to be neurotoxic in striatal neurons (Ben-Shachar et al. 1995).

#### 2.2 Neurotoxicity Induced by Calcium Ions

High  $Ca^{2+}$  cell concentration in the cytosol initiates apoptosis (a cell death that leads to the elimination of cells without releasing harmful substances into the surrounding area) through cleaved caspase processing and activation of a number of several intracellular enzymes (e.g., phospholipase A<sub>2</sub>, nitric oxide synthase, calcineurin, endonucleases, and xanthine dehydrogenase) and proteases (e.g., calpain) which by themselves can induce the formation of RS or damage the cytoskeleton, membrane, or DNA (Rego and Oliveira 2003). Another damaging result of excess Ca<sup>2+</sup> in the cytosol is the opening of the mitochondrial permeability transition pore. Opening this mitochondrial pore may cause release of reactive oxygen species (ROS), more Ca<sup>2+</sup>, more proteins that can lead to apoptosis and finally to cessation of adenosine triphosphate (ATP) production (Court and Coleman 2012).

A rise in  $Ca^{2+}$  intracellular concentration is generated – among others – by glutamate high synaptic concentrations (see above) or products of lipid peroxidation (e.g., 4-hydroxynoneral; see below).

## 2.3 Neurotoxicity Induced by RS

RS include free radicals and non-radical molecules. A free radical means any species capable of independent existence ("free") that contains one or more unpaired valence shell electrons (Halliwell and Gutteridge 1988). These chemically reactive molecules include ROS, reactive bromine species (RBS), reactive chloride species (RCS), and reactive nitrogen species (RNS). It should be stated that the term ROS is usually used as a collective term for both oxygen radicals that include free radicals (e.g., superoxide O2., hydroxyl OH, hydroperoxyl HO2) and nonradicals (e.g., hydrogen peroxide H<sub>2</sub>O<sub>2</sub>, peroxynitrite ONOO<sup>-</sup>). The collective definition exists also for RBS (free radical: atomic bromide Br and non-radicals: bromine gas Br<sub>2</sub>, hypobromous acid, HOBr), for RCS (free radical: atomic chlorine Cl and non-radicals: hypochlorous acid HOCl, chloramines), and for RNS (free radicals: nitric oxide NO, nitrogen dioxide NO<sub>2</sub>, and nitrate NO<sub>3</sub> and non-radicals: nitrous acid HNO<sub>2</sub>, nitrosyl anion NO<sup>-</sup>, peroxynitrite ONOO<sup>-</sup>, and peroxynitrate  $O_2NOO^-$ ). Among RS, the most neurotoxic are  $O_2^{\bullet-}$ ,  $OH^{\bullet}$ , and  $H_2O_2$  (Bartosz 2009: Matés et al. 2010), the first radical leaking from the respiratory chain in mitochondria, the second radical resulting from the dismutation of  $O_2^{\bullet-}$  or directly from the action of oxidase enzymes, and the non-radical species causing modification of purine and pyrimidine bases that lead to DNA damage.

The most important source of RS for aerobic cells is O<sub>2</sub> due to its high consumption, especially by the mammalian brain (Halliwell 2006). When formed during the normal metabolism of oxygen, RS have important roles in cell signaling and homeostasis; however, strong environmental stressors (e.g., UV, heat exposure, ionizing radiation, chemical substances including addictive drugs) increase their brain levels. In the brain, formation of oxygen radicals is linked to an increase in the  $Ca^{2+}$  concentration inside a cell (via membrane  $Ca^{2+}$  exporters, transient receptor potential melastatin-related (TRPM)2 cation channels located on some neurons and glia or via Ca<sup>2+</sup> pumps in the endoplasmic reticulum) due to damage of mitochondrial function by neurotoxins following disruption of ATP production. High intracellular Ca<sup>2+</sup> levels can activate neuronal nitric oxide synthase (NOS), phospholipase A<sub>2</sub>, and calpains and/or interfere with mitochondrial function. The interaction between the Ca<sup>2+</sup> and Ca<sup>2+</sup>-dependent NOS pathway results in the production of NO<sup>•</sup> which reacts with O<sub>2</sub> forming ONOO<sup>-</sup>. ONOO<sup>-</sup> and other RS may decrease glutamate transporter activity on glial cells and inactivate glutamine synthethase (Aksenov et al. 1997; Trotti et al. 1996; Noack et al. 1998; Koppal et al. 1999), while 4-hydroxynonenal (a major aldehydic product formed by peroxidation of  $\omega$ -6-unsaturated fatty acids regarded as a specific marker of lipid peroxidation) can impair the function of glutamate transporters (Mattson and Chan 2003). Disturbances in glutamate clearance may promote the high Ca<sup>2+</sup> intracellular levels and lead to mitochondrial and protein dysfunctions.

Other potential sources of RS are cytochromes P450 (CYPs) and activated microglia. CYPs are enzymes localized to almost every cell, including some brain regions (Miksys and Tyndale 2004). Among CYPs, the CYP2E1 leaks electrons during the catalytic cycle and is especially linked to RS production (Gonzalez 2005);

however, the CYP2E1-dependent RS generation is important following exposure to, e.g., solvents, tobacco, or ethanol, as these substances significantly increase the enzyme levels.

It was also found that neurotoxin-activated microglia can produce  $O_2^{\bullet-}$ ,  $H_2O_2$ , and cytokines (interleukin (IL)-1, interleukin-6, and TNF  $\alpha$ ). In turn, cytokines lead to activation of microglia and then to production of more RS, inducible NOS and NO<sup>•</sup> (Duncan and Heales 2005). Of note, cytokines may be also produce by activated astroglia (Duncan and Heales 2005).

RS are also generated by endogenous sources such as neurotransmitters. In fact, catecholamines such as dopamine, its precursor L-DOPA and norepinephrine, as well as indolamine serotonin (5-HT) react with  $O_2$  to produce  $O_2^{\bullet-}$  (Spencer et al. 1998). The best recognized is dopamine-induced generation of RS. Thus, dopamine metabolism includes either a nonenzymatic pathway (via molecular oxygen through autoxidation) to generate hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) and a superoxide anion (O<sub>2</sub><sup>•-</sup>) or an enzymatic pathway by monoamine oxidase (MAO) to form H<sub>2</sub>O<sub>2</sub>. Both of these radicals may react via the Haber-Weiss/Fenton reaction with transition metal ions to produce the highly toxic OH<sup>•</sup> (Hermida-Ameijeiras et al. 2004).

As the brain is the most vulnerable to increased levels of RS due to its low oxidative capacity and because neurons are characterized as postmitotic cells, it means that neurons live with accumulated damage over the years. Harmful effects of RS on the cell include induction of apoptotic pathways, damage of DNA, oxidations of polyunsaturated fatty acids in lipids (lipid peroxidation and malondialdehyde (MDA) are widely studied; MDA is a product of polyunsaturated fatty acid peroxidation), oxidations of amino acids in proteins, and oxidatively inactivated specific enzymes by oxidation of their cofactors. Of note, there is positive feedback response between the mitochondrial production of RS and glutamate concentration at the synapse which further gives rise to postsynaptic Ca<sup>2+</sup> levels via RS-mediated inhibition of glial EAAT2 function.

Whether RS will act as damaging or signaling molecules and whether "oxidative stress" occurs depends on the delicate equilibrium between RS production and scavenging/detoxification. There are several cellular or extracellular defensive mechanisms comprising of the nonenzymic as well as enzymic antioxidants to break down RS. The enzymic antioxidants include proteins such as superoxide dismutases (SOD), catalases (CAT), peroxiredoxins, glutathione peroxidase (GPx), glutathione reductases (GR), thioredoxin reductases, heme oxygenase-1, eosinophil peroxidase, and metallothionein (Matés et al. 2010). An essential defense role against oxidative stress in all aerobic organisms is played by SOD which converts  $O_2^-$  to  $H_2O_2$  and molecular oxygen. Four classes of SOD have been identified: cytosolic Cu, Zn-SOD, mitochondrial Mn-SOD, extracellular-SOD, and mitochondrial Fe-SODs; the first three SODs exist in humans. CAT, GRx, or peroxiredoxins convert molecules of  $H_2O_2$  into water and oxygen, while GR catalyzes the reduction of glutathione disulfide (GSSG) to glutathione (GSH) and, thus, maintains a cellular redox state (Matés et al. 2008).

A potent nonenzymatic antioxidant defense system is comprised of vitamin C (ascorbic acid), vitamin E ( $\alpha$ -tocopherol), vitamin A,  $\beta$ -carotene, polyamines,

melatonin, NADPH, adenosine, urate, coenzyme Q-10 (ubiquinol), polyphenols, flavonoids, phytoestrogens, GSH, cysteine, homocysteine, taurine, methionine, *S*-adenosyl-L-methionine, nitroxides, and selenium (Matés et al. 2008). Among these small antioxidant molecules ascorbate, GSH and tocopherols are of special significance. Thus, ascorbic acid donates electrons in a number of enzymatic and nonenzymatic reactions, while GSH by reaction with OH<sup>•</sup> or  $H_2O_2$  can function directly as a free radical scavenger and protect proteins, lipids, or DNA either by the formation of adducts directly with reactive electrophiles or by acting as proton donor in the presence of RS. Tocopherols potently scavenge oxygen and protect lipids and lipid structures against peroxidation.

## 2.4 Neurotoxicity Induced by Apoptosis

Due to physiological as well as pathological conditions including intracellular rise in RS or  $Ca^{2+}$  levels as well as other strong stressors (e.g., hyperthermia, ionizing radiation, some toxic substances; (Bojes et al. 1999; Darzynkiewicz et al. 2009)), a programmed cell death (apoptosis) appears. Apoptosis is a multistep process characterized by biochemical events that lead to cell blebbing, cell shrinkage, nuclear fragmentation, chromatin condensation, chromosomal DNA fragmentation, and, finally, cell death. The process of apoptosis is controlled by diverse signals, which may originate either extracellularly (e.g., toxins, cytokines) that must cross the plasma membrane or transduce a response or intracellularly (Darzynkiewicz et al. 2009).

Apoptosis is executed by either caspases or noncaspase proteases (Wyllie 2010). Caspases are cysteine-dependent aspartate-specific proteases forming a family of over a dozen caspases. They function as initiator caspases (caspase 2, 8, 9, 10 leading to activation of effector caspases) and effector caspases (caspase 3, 6, 7 that proteolytically degrade intracellular proteins to kill a cell). Two intracellular caspase pathways have been recognized; one is triggered by message from a death receptor (so-called the extrinsic pathway) and the other activated by an intracellular message such as DNA damage and microtubule disruption (so-called the intrinsic or mitochondrial pathway) (Gabai et al. 2002).

The extrinsic pathway for caspase activation engages apoptotic mechanisms through stimulation of the TNR family, i.e., TNF receptors and Fas at the cell surface. TNF1 receptors are activated by TNF, and such linkage results in the release of silencer of death domains (SODD) and formation of a receptor-proximal complex containing the important adaptor proteins TNF receptor-associated death domain (TRADD), TNF-R-associated factor 2 (TRAF2), receptor-interacting protein (RIP), and Fas-associated death domain (FADD). These adaptor proteins enhance activity of the TNF pathway-specific enzymes (e.g., caspase 8 and IKK $\beta$ ) to the TNF-R1 complex. These enzymes, in turn, initiate the activation of c-Jun NH<sub>2</sub>-terminal kinase (JNK), a kinase that phosphorylates c-Jun and increases its transcriptional activity, while the protein kinase RIP is critical to the activation of the transcription factor NF- $\kappa$ B.

Fas are also stimulated by another death activator, Fas ligand (FasL). The interaction between Fas and FasL results in the formation of the death-inducing signaling complex (DISC), which contains the FADD, caspase 8 and caspase 10. Activation of caspase 8 initiates a cascade of caspase activation leading to cell phagocytosis.

The internal/mitochondrial pathway for caspase activation is initiated by proapoptotic (Bax, Bak, Bid, or Bad) homodimers of the *Bcl-2* family that inhibit antiapoptotic (Bcl-Xl and Bcl-2) Bcl-2 members. The proapoptotic proteins are required to allow the release of caspase activators such as cytochrome c and small mitochondria-derived activator of caspases (SMAC) out the mitochondrial membrane. The released cytochrome c binds to the apoptotic protease activating factor-1 (Apaf-1) forming apoptosomes that later activate caspase 9. Caspase 9, in turn, activates effector caspases 3 and 7. Caspase 3 destroys poly-ADP ribose polymerase (PARP), a protein which takes part in DNA repair; caspase 3 also initiates DNA fragmentation. SMACs are mitochondrial proteins released into the cytosol to bind to the inhibitor of apoptosis proteins (IAPs) such as caspases and deactivate them.

The noncaspase-related execution of apoptosis is linked to activation of calpains. The calpains exist in two isoforms (I and II) and are  $Ca^{2+}$ -dependent proteases. They degrade membrane, cytoplasmic, and nuclear substrates, leading to the breakdown of cellular architecture. Among others, they release proapoptotic factor (AIF) (Breckenridge and Xue 2004) from the mitochondria into the cytosol. AIF migration into the nucleus is mediated by its nuclear localization signal (NSL); in the nucleus AIF binds with DNA strands inducing its fragmentation and cell death. Of note, another effector of caspase-independent cell death is endonuclease G (Endo-G) released from mitochondria (Singh et al. 2004).

#### 2.5 Neurotoxicity Induced by Glial Cell Activation

Among several glial cells, microglia are preferentially exposed to toxic factors and are strongly engaged in brain toxicity.

Microglia constitute 20 % of the total glial cell population within the brain and are the first form of active immune defense in the CNS responsible for maintaining homeostasis. There are several factors that activate microglia, such as glutamate and glutamate receptor agonists, proinflammatory cytokines, cell necrosis factors, lipopolysaccharide, and extracellular potassium changes (Nakajima and Kohsaka 2001; Graeber and Streit 2010). In turn, activated microglia via hemichannels within cell membranes (Orellana et al. 2012) can secrete a variety of neurotoxic substances, among them are (i) large amounts of  $H_2O_2$  and NO<sup>•</sup> – the process known as "respiratory burst" leads to neuronal cell death; (ii) proteases (i.e., cathepsins B, L, and S; the matrix metalloproteinases (MMP)-1, MMP-2, MMP-3, and MMP-9; the metalloprotease-disintegrin ADAM8 plasminogen; and elastase) that catabolize various proteins causing cellular damage; (iii) nuclear factor  $\kappa$ B, a transcription factor that starts proinflammatory actions in microglia; (iv) proinflammatory chemokines (MCP-1, MIP-1 $\alpha$ , and MIP-1 $\beta$ ); (v) proinflammatory cytokines (IL-1 $\alpha$ , IL-1 $\beta$ ) and TNF- $\alpha$  that initiate demyelination of neuronal axons and/or apoptosis through activation of caspase 8 and caspase 3; and (vi) excitatory amino-acid glutamate and aspartate that injury neurons (Block et al. 2007; Graeber and Streit 2010).

Another glial cell type, oligodendroglia, expresses a variety of ligand- and voltage-gated neurotransmitter receptors/ion channels, including NMDA and AMPA (Belachew et al. 1999; Rogister et al. 1999). The presence of functional neurotransmitter receptors/ion channels in glia indicates that there is glutamate-mediated cross talk between neurons and glia (Bergles et al. 2000; Gallo and Ghiani 2000) as well as that which oligodendrocytes share with neurons – a high vulnerability to glutamate excitotoxicity.

On the other hand, oligodendrocytes (as well as astrocytes) have the ability to protect themselves from RS as they contain high concentrations of antioxidant enzymes and GSH (Makar et al. 1994). However, in conditions where protective functions of astrocytes are impaired, oligodendrocytes may be vulnerable to oxidative and RS-mediated damage (e.g., following increased metabolism of catecholamine neurotransmitters that generates  $H_2O_2$ ) (Noble et al. 1994).

## 2.6 Neurotoxicity Induced by DNA Damage

DNA is a genetic material, and any DNA damage can result in malfunctions or complete inactivation of encoded proteins. Genomic instability is an important mechanism leading to death of neurons. Several environmental factors (e.g., UV radiation, plant and chemical toxic substances, viruses) or endogenous sources (e.g., RS) are responsible for DNA damage that may occur during the synthesis of DNA precursors, DNA repair, and during various methylation events, including DNA methylation (see below). Both DNA repair and DNA methylation are critical for maintaining genomic stability. Several forms of DNA damage may appear including mutations, deletions, gene amplification, and rearrangements (Matés et al. 2008).

There is a relationship between oxidative stress and genomic instability (Matés et al. 2008). In fact, ROS (especially OH<sup>•</sup>) are a major source of DNA sequence changes leading to deoxyribose oxidation, a generation of a deoxyribose radical that further initiates strand breakage, removal of nucleotides, a variety of modifications in the organic bases of the nucleotides, and DNA-protein cross-linkages. Of note, the formation of 8-hydroxy-2'-deoxyguanosine is regarded as an index of oxidative DNA damage (Frenkel et al. 1991).

DNA damage may result in the initiation of signaling leading to neurotoxic consequences such as cell death or to the activation of several proto-oncogenes and/ or the inactivation of some tumor suppressor genes such as p53 (Hetman et al. 2010; Raj et al. 2011). p53 gene is a suppressive gene coding a p53 protein – a tumor suppressor protein and its level in unstressed cells are kept low through a continuous degradation, but DNA damage and other stress signals trigger the increase of p53 protein levels and its activity. The major functions of p53 protein are (i) a growth arrest (a stop signal for progression of cell cycle) during DNA damage and

(ii) induction of DNA repair, and when unsuccessful, it directs cells on the road to apoptosis via an upregulation of Bax, Fas, and Apaf- 1 expression (Hetman et al. 2010; Raj et al. 2011).

## 2.7 Neurotoxicity Induced by Epigenetic Mechanisms

Apart from genetic material, epigenetic regulation of gene expression (=epigenetic processes) can change dynamically in response to external factors, providing a key mechanism by which the environment can influence gene expression and, hence, phenotype (Taft et al. 2007; Archer et al. 2010). Epigenetic mechanisms reflect the regulation of gene expression, occurring independently of DNA sequence (Henikoff and Matzke 1997). There were recognized – at least – three forms of epigenetic mechanism: DNA methylation, histone tail modifications, and regulatory noncoding RNA (ncRNA).

DNA methylation occurs primarily at cytosine-guanine dinucleotides (CpG) and requires DNA methyltransferases (DNMTs). The addition of a methyl group to CpG sites displaces the binding of transcription factors and attracts methyl-binding proteins that instigate chromatin compaction of DNA and gene silencing.

Histones are the basic proteins around which DNA is wrapped to form nucleosome. Covalent histone tail modifications, including methylation, acetylation, phosphorylation, sumoylation, and ubiquitylation, modulate gene expression via alterations in chromatin structure. Acetylation, one of the most frequent posttranscriptional modifications of histones, depends on histone-modifying enzymes, such as histone acetyltransferases (HATs) and histone deacetylases (HDACs). Generally, histone acetylation belongs to the active chromatin marks; it increases the accessibility of transcription complexes to genomic DNA. Histone methylation, controlled by several families of histone methyltransferases (HMTs), has been associated with both transcriptional activation and repression, depending on the particular residue and the extent of methylation (Su and Tarakhovsky 2006; Maze and Nestler 2011); this reaction can be reversed by equally diverse histone demethylases (HDMs).

Regulatory ncRNAs (encoded by part of DNA noncoding proteins) are able to control transcription and translation in an organ- and cell-specific manner; the disturbance of their function could lead to perturbation in cellular homeostasis. These include, inter alia, microRNAs (miRNAs), Piwi-interacting RNAs (piRNAs), endogenous small interfering RNAs (endo-siRNAs, esiRNAs), and long noncoding RNAs (lncRNAs).

miRNAs represent a class of short (~18–25 nucleotides in length), single-stranded molecules which cause epigenetic changes in gene transcription, mediated by both DNA methylation and histone modifications (Kim et al. 2007, 2009). miRNA are expressed in the mature mammalian brain (Landgraf et al. 2007) and are thought to be involved in long-term potentiation and in regulating structural and functional aspects of synaptic plasticity (i.e., neuronal morphogenesis, activity-dependent translation during synapse formation, memory and synapse development, and plasticity (Schratt 2009)). miRNAs in the vertebrate system predominantly inhibit productive translation

of mRNAs in the cytoplasm (Schratt et al. 2006) and may regulate transcriptomic networks via the expression of other types of lncRNAs in the nucleus (Zhou et al. 2010). On the other hand, the expression of miRNAs is modulated by DNA methylation and histone modifications.

piRNAs are a class of small noncoding RNA, 24–31 nucleotides in length that forms RNA-protein complexes with Piwi proteins (Dharap et al. 2011). piRNAs are present in the polysomal fraction, but surprisingly they were also found in the nucleus. piRNAs play an important role in silencing pathways that regulate mRNA stability, heterochromatin formation, and genome integrity (Brower-Toland et al. 2007). They may guide DNA methylation and control transposon activity (Xu et al. 2008; Zhou et al. 2010), and in turn, impaired epigenetic regulation of transposons can interfere with the expression of neighboring genes. Recent studies confirmed their expression in somatic tissues (Spadaro and Bredy 2012) and in the epigenetic facilitation (chromatin remodeling) of long-term memory and neuronal plasticity (Lee et al. 2011; Rajasethupathy et al. 2012; Spadaro and Bredy 2012).

esiRNAs represent one of several classes of small noncoding RNAs involved in RNA interference. esiRNA is derived from repeated DNA elements and transposons (Obbard and Finnegan 2008). In addition to their role as posttranscriptional regulators, esiRNAs function as epigenetic regulators through either RNA-directed DNA methylation or RNA-mediated heterochromatin formation (Verdel et al. 2009; Lejeune and Allshire 2011). In both cases, esiRNAs may act as guide molecules to direct epigenetic factors such as site-specific MNMT1 or HMTs to target specific genomic regions (Song et al. 2011). Chen et al. (2012) have demonstrated that esiRNAs by silencing LINE-1 activity through DNA methylation in human cells can function as a link between the expression of esiRNAs and LINE-1 retrotransposons. The recent study has also showed that adult mouse hippocampus expresses esiRNAs that possibly play more specific roles in regulating synaptic plasticity and learning (Smalheiser et al. 2011).

lncRNAs are mRNA-like transcripts ranging in length from 200 nucleotides to  $\sim$ 100 kilobases that do not have functional open reading frames. Most lncRNAs are localized in nucleus, display low expression compared to protein coding genes, and are composed of both polyA+ and polyA- transcripts (Kaikkonen et al. 2011). lncRNA can be divided into five categories based on their proximity to coding genes: intronic, intragenic, bidirectional, sense, and antisense (Qureshi and Mehler 2011). They have been associated with various cellular processes, such as transcription regulation (in cis or trans loci), alternative splicing, control of the subcellular localization of transcription regulatory factors and proteins, chromosomal dosage compensation, as well as chromatin remodeling (Johnson et al. 2012; Li et al. 2012). IncRNAs recruit histone modification complex to target sites in chromatin and thus can function as epigenetic regulatory factors (Rinn and Chang 2012). They exhibit cell-specific patterns of expression within brain, suggesting their role in the modulation of genes associated with neuronal connectivity and synaptic plasticity (Bernard et al. 2010; Li et al. 2012). Recently, IncRNAs were shown as factors engaged in GABA neuron neurogenesis (Mercer et al. 2010).

## 3 Conclusion

Among endogenous brain toxins, high concentrations of glutamate, calcium ions, and RS are well-known factors that potently disturb the normal physiology and function of the nervous system. It was also established that glial cells (i.e., astroglia, microglia, and oligodendrocytes) and their released factors (e.g., proinflammatory cytokines and chemokines, excitatory amino acids, cell necrosis factors, RS) also play a number of active roles in the brain and are sensitive indicators for pathology of the CNS. Recently, genomic instability due to DNA damage and epigenetic mechanisms (such as DNA methylation, histone tail modifications, and regulatory ncRNAs) were recognized as important mechanisms leading to disturbances of neurons and finally to their death.

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# Quinolinate and Related Excitotoxins: Mechanisms of Neurotoxicity and Disease Relevance

# Jonas I. Addae and Trevor W. Stone

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J.I. Addae (🖂)

Department of Preclinical Sciences, University of the West Indies, St Augustine, Trinidad and Tobago e-mail: jonas.addae@sta.uwi.edu

T.W. Stone

Institute of Neuroscience and Psychology, University of Glasgow, Glasgow, UK e-mail: trevor.stone@glasgow.ac.uk

#### Abstract

There are many ways in which neuronal damage can be produced in the brain, including the overactivation of depolarizing receptors, exposure to high levels of pro-inflammatory proteins such as cytokines, or miscellaneous toxins, but the kynurenine pathway has emerged as a novel but potentially major factor in regulating neuronal viability or death. It is the major route for the metabolism of the essential amino acid tryptophan, which is oxidized by indoleamine-2,3-dioxygenase (IDO) to a series of compounds which can activate, block, or modulate conventional neurotransmitter receptors. Quinolinic acid is an agonist at N-methyl-D-aspartate receptors, kynurenic acid is an antagonist at these and other glutamate receptors, and other kynurenine metabolites are highly redoxactive. Superimposed on the discovery of this neuromodulatory pathway have been observations that activity in the pathway is linked to neurological and psychiatric disorders, correlating with disease state (as in Huntington's disease) or cognitive function (as following bypass surgery). Together, the data accumulated to date make a strong case for this hitherto obscure pathway being a major factor in determining cell damage, death, or recovery in health and disease.

#### **Keywords**

Alzheimer's disease • Depression • Excitotoxicity • HIV-1-Associated neurocognitive disorder • Huntington's disease • Kynurenate • Kynurenine pathway • Multiple sclerosis • Neurodegeneration • Neuro-inflammation • Parkinson's disease • Quinolinic acid • Schizophrenia • Stroke

LISC OF AD	bicviations
3HA	3-Hydroxyanthranilic acid (3HA)
3HAO	3-Hydroxyanthranilic acid oxygenase
3HK	3-Hydroxkynurenine
AD	Alzheimer's disease
AIDS	Acquired immunodeficiency syndrome
AMPA	α-Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid
Gly	Glycine
HD	Huntington's disease
HIV	Human immunodeficiency virus
IDO	Indole amine 2,3-dioxygenase
IFN	Interferon
KAT	Kynurenine amino transferases
LPS	Lipopolysaccharides
MAPK	Mitogen-activated protein kinase
mGluR	Metabotropic glutamate receptor
MPTP	1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine
MS	Multiple sclerosis
nAChR	Nicotinic acetylcholine receptor
$NAD^+$	Nicotinamide adenine dinucleotide

NMDA	<i>N</i> -methyl-D-aspartate
PD	Parkinson's disease
TDO	Tryptophan 2,3-dioxygenase
TNF	Tumor necrosis factor

#### 1 Introduction

Quinolinic acid is a highly selective agonist at the population of glutamate receptors that respond to the synthetic amino acid *N*-methyl-D-aspartate (NMDA). In contrast to NMDA, however, quinolinic acid is an endogenous compound, generated via the kynurenine pathway for the oxidation of another amino acid, tryptophan (Stone and Perkins 1981). The discovery that quinolinic acid is able to produce depolarization by activating NMDA receptors (NMDARs) led to the demonstration that it also causes neurodegeneration (Schwarcz et al. 1983), consistent with the knowledge that glutamate itself can induce neuronal damage by promoting depolarization and an excessive influx of calcium. These discoveries launched the concept that the brain damage associated with stroke or degenerative disease might be attributable not only to the accumulation of glutamate or hyperactivity of its receptors, but also to abnormal function of the kynurenine pathway.

The kynurenine pathway also produces redox-active compounds (3-hydroxykynurenine and 3-hydroxyanthranilic acid) which cause neuronal damage by generating reactive oxygen species (Eastman and Guilarte 1989; Giles et al. 2003). A branch of the pathway generates kynurenate which is an antagonist of glutamate (Perkins and Stone 1982) and nicotinic cholinergic receptors (Hilmas et al. 2001; Stone 2007). As a result, kynurenate exhibits neuroprotective properties. A number of diseases have been associated with altered levels of quinolinic acid and other kynurenines in the CNS, including the AIDS-dementia complex, Alzheimer's disease, Huntington's disease, and Parkinson's disease. In addition, as the relevance of neuronal viability and damage become clear in the etiology of psychiatric disorders such as schizophrenia and depression, and inflammatory diseases such as multiple sclerosis, the pathological and therapeutic importance of kynurenine metabolism continues to expand, as indicated in the sections that follow.

## 2 Components of the Kynurenine Pathway

#### 2.1 Tryptophan Cleavage and Immune System

L-tryptophan, the least abundant of the essential amino acids used for protein synthesis, is also a substrate for synthesizing the neurotransmitter serotonin via the methoxyindole pathway, or vitamin B3 (nicotinamide) and quinolinic acid via the kynurenine pathway (Fig. 1).

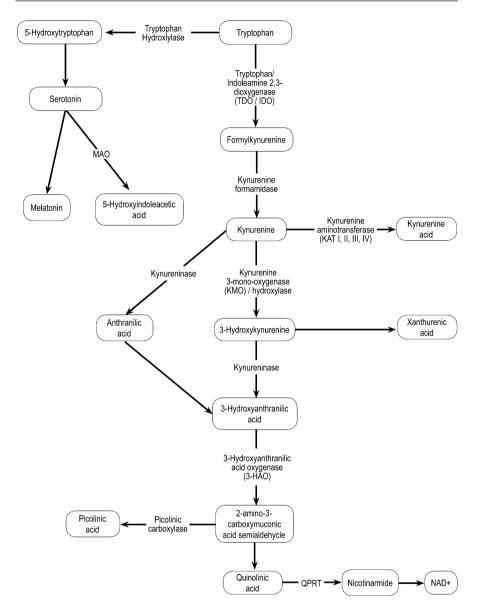


Fig. 1 A summary of the major compounds in the kynurenine pathway of tryptophan oxidation

Most of the ingested tryptophan (over 95 %) is catabolized in the kynurenine pathway (Stone 1993a; Takikawa 2007) and generates nicotinamide adenine dinucleotide (NAD<sup>+</sup>) under physiological conditions. L-tryptophan is metabolized to kynurenine by oxidative cleavage of the indole ring by the enzymes tryptophan 2,3-dioxygenase (TDO) or indoleamine 2,3-dioxygenase (IDO), the activities of which may act as rate-limiting steps under some conditions (Takikawa et al. 1986;

Rafice et al. 2009). IDO, a heme-containing inducible enzyme, is found in the lungs, blood, and brain (Heyes et al. 1993; Banks et al. 1994) and has a higher affinity for tryptophan (Km =  $\sim 10 \mu$ M) than does TDO (Km =  $\sim 200 \mu$ M). IDO has recently been shown to exist in at least two isoforms, though little is yet known about IDO-2 (Ball et al. 2009).

A key aspect of the kynurenine pathway that is becoming increasingly recognized for its importance in neurodegeneration is its intimate relationship with the immune system. The IDO promoter has one interferon- $\gamma$  (IFN- $\gamma$ ) activation site and two nonspecific IFN-stimulated response elements that respond to IFN- $\alpha$ , IFN- $\beta$ , or IFN- $\gamma$ , with IFN- $\gamma$  being the most potent inducer (Puccetti 2007). Pro-inflammatory interleukins such as IL-10 activate T cells and natural killer cells to produce IFN- $\gamma$ which acts synergistically with tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) to increase expression of IDO (Robinson et al. 2003). During immune system activation, this sequence serves as a form of defense against infection by causing tryptophan depletion (Takikawa 2007). IDO can be activated also by some pro-inflammatory cytokines (IL-18, IL-12) independently of IFN- $\gamma$  (Liebau et al. 2002).

In the CNS, IDO is found in microglia, macrophages, astrocytes, neurons, and microvascular endothelial cells (Guillemin et al. 2001, 2005, 2007). Activated microglia produce pro-inflammatory cytokines IL-1 $\beta$ b, IL-2, IL-6, TNF- $\alpha$ , and IFN- $\gamma$  which cause further microglial activation to amplify the inflammatory response.

TDO is found mainly in the liver but is also present in several brain areas including the cerebral cortex, cerebellum, hypothalamus, and brainstem (Haber et al. 1993; Kanai et al. 2009). The enzyme is readily activated by IFN- $\gamma$  or bacterial endotoxins (Yoshida et al. 1981).

# 2.2 Formation of 3-Hydroxy-Kynurenine, 3-Hydroxy-Anthranilic Acid, and Quinolinic Acid

Kynurenine is transported easily into the CNS across the blood–brain barrier by the large neutral amino acid transporter; hence, changes in the plasma level of kynurenine are reflected in CSF levels (Fukui et al. 1991; Jauch et al. 1993). Approximately 60 % of the brain kynurenine is from the systemic circulation (Gal and Sherman 1978). Brain immune cells take up kynurenine from the extracellular space via the Na<sup>+</sup>-independent large neutral amino acid transporter (Speciale et al. 1989).

Kynurenine 3-monoxygenase (KMO) [also known as kynurenine 3-hydroxylase], which uses kynurenine as a substrate, is present mainly in the liver. It is also present in the brain albeit with a much lower activity than in peripheral tissues. KMO is located in the outer membrane of mitochondria and is not saturated under physiological conditions. The production of 3-hydroxkynurenine (3HK) can, therefore, be increased when there is increased supply of kynurenine, as, for example, from the periphery (Schwarcz et al. 2010). The cytokine IFN- $\gamma$  activates KMO; hence, inflammatory states stimulate the expression of KMO and production of quinolinic acid (Heyes and Morrison 1997; Connor et al. 2008). In the CNS, resident microglia and invading macrophages from the periphery express far more KMO and 3-hydroxyanthranilic acid oxygenase (3HAO) enzymes than astrocytes, and therefore are the major source of quinolinic acid (Heyes et al. 1996; Lehrmann et al. 2001).

#### 2.3 Formation of Kynurenate

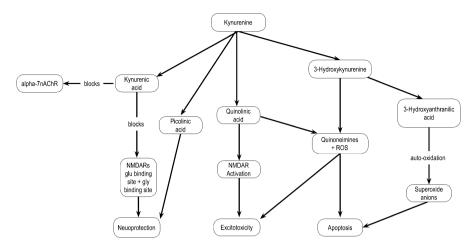
In mammalian brain, kynurenine amino transferases (KAT), the enzymes that convert kynurenine to kynurenate, are found mainly not only in astrocytes but also in neurons to some extent (Guillemin et al. 2000; Guidetti et al. 2007b). Kynurenate is produced by the irreversible transamination of kynurenine by four mammalian pyridoxal-5'-phosphate dependent enzymes – KAT I (glutamine transaminase or cysteine conjugate beta-lyase I), KAT II (alpha aminoadipate aminotransferase), KAT III (cysteine conjugate beta-lyase II), and KAT IV (glutamic-oxaloacetic transaminase or mitochondrial aspartate aminotransferase). KAT II accounts for most (50–60 %) of kynurenate formation in human and rat brains, whereas KAT IV, a more recent addition, is the dominant enzyme (60 %) in mouse brain (Guidetti et al. 2007a; Han et al. 2010).

Once formed, kynurenate cannot be metabolized further and is released into the extracellular compartment by a non-Ca<sup>2+</sup> -dependent process (Turski et al. 1989; Kiss et al. 2003). Kynurenate is removed slowly from the brain by the probenecid-sensitive, nonspecific acid transporter and by passive diffusion (Moroni et al. 1988; Schwarcz and Pellicciari 2002). Although KMO is in the pathway that produces 3HK and quinolinic acid, its activity regulates the levels of kynurenine in the brain, and, hence, indirectly affects the rate of formation of kynurenate (Wonodi and Schwarcz 2010).

#### 3 Neurodegeneration and Kynurenine Metabolites

Excitotoxicity was first described by Olney et al. (1975) as the damage to neurons that results from excessive stimulation of glutamate receptors (Olney et al. 1975). The neuronal damage caused by stimulation of glutamate receptors does not affect the axons of passage or the afferent nerve terminals. Endogenously released glutamate does not cause much excitotoxicity due to the very efficient uptake systems (Danbolt 2001); hence, quinolinic acid, being an endogenous excitotoxin with no efficient removal mechanism, has been of particular interest for several years.

The concentration of quinolinic acid in the brain is normally too low to cause acute toxicity although maintained high nanomolar concentrations may damage or kill particular groups of neurons (Giulian et al. 1990). During inflammation, quinolinic acid may increase from the normal nM levels to micromolar levels which can cause overt and rapid neurotoxicity (Heyes et al. 1998). Nonetheless, quinolinic acid has been reported to be toxic to neurons even at concentrations as low as 150 nM (Braidy et al. 2009).



**Fig. 2** A diagrammatic representation of how the role of the kynurenine pathway in regulating neuronal death or survival depends critically on the ratio between compounds such as quinolinic acid, which can be neurotoxic in excessive concentrations, and kynurenic acid, which is protective by blocking NMDA and nicotinic receptors. The contributions of redox-active compounds such as 3-hydroxykynurenine and 3-hydroxyanthranilic acid depend on their relative concentrations as well as the ambient redox state of the tissue

Quinolinic acid produced by microglia and infiltrating macrophages acts on NMDARs (Addae and Stone 1987; Stone 1993a), particularly those with NR1/NR2A or NR1/NR2B subunits (Perkins and Stone 1983; Stone 1993b; de Carvalho et al. 1996), and causes neurotoxicity partly via overactivation of these receptors (Fig. 2). Excessive stimulation of NMDARs causes influx of Ca<sup>2+</sup> which overcomes cellular Ca<sup>2+</sup> sequestration processes and leads to activation of proteases, endonucleases, generation of reactive oxygen species and nitric oxide that eventually cause neuronal death by lipid peroxidation or apoptosis. In addition, quinolinic acid stimulates hypertrophy and proliferation of astrocytes which then release increased amount of cytokines, leading to astrogliosis and inhibition of glutamate reuptake to cause further excitation of NMDARs (Tavares et al. 2002; Ting et al. 2009).

Two other products in the kynurenine pathway, 3-hydroxykynurenine (3HK) and 3-hydroxyanthranilic acid (3HA), cause neurodegeneration by generating superoxide and hydrogen peroxide under appropriate redox conditions (Goldstein et al. 2000). The axon-sparing excitotoxicity of quinolinic acid is potentiated by the presence of 3HK and reactive oxygen species; this synergistic effect is unique to quinolinic acid, because there is no such effect between NMDA and 3HK (Guidetti and Schwarcz 1999; Behan and Stone 2002). This synergistic effect explains why, although the potency of quinolinic acid that causes damage in the CNS is about the same as that of NMDA.

The activity of 3HAO, the oxygenase enzyme that converts 3-hydroxyanthranilic acid to quinolinic acid, is iron  $(Fe^{2+})$  dependent. Hence,

excessive iron levels in the brain dramatically increase the levels of quinolinic acid which may contribute to the neurodegeneration caused by iron (Stachowski and Schwarcz 2011). Quinolinic acid itself may also cause neurotoxicity via formation of iron complexes, leading to lipid peroxidation (Platenik et al. 2001).

# 4 Neuroprotective Kynurenine Metabolites and Their Control

Kynurenate is the only known endogenous antagonist of NMDARs that preferentially blocks the glycine co-agonist binding site (Gly<sub>2</sub> or Gly<sub>B</sub>) where it has a higher affinity for the glycine site than glutamate (Birch et al. 1988; Kessler et al. 1989). Kynurenate is normally present in the brain at average levels that are in the high nanomolar range (200–1,500 nM) (Turski et al. 1988), but at higher micromolar concentrations, which are likely to exist near the sites of production and cellular efflux, kynurenate blocks competitively all ionotropic glutamate receptors (Perkins and Stone 1982; Stone 1993a). This effect of kynurenate on non-NMDA glutamate receptors at synaptic sites enables it to effectively block thalamic projections to neocortical neurons (Addae and Stone 1988). Kynurenate also blocks noncompetitively the  $\alpha$ 7-nicotinic acetylcholine receptor (nAChR), and inhibits release of glutamate from presynaptic terminals by acting on this type of receptor (Hilmas et al. 2001; Carpenedo et al. 2001).

One consequence of the antagonistic activity of kynurenate at glutamate receptors is that a reduction in the levels of brain kynurenate promotes NMDAR-mediated excitotoxicity (Sapko et al. 2006). Such a reduction may occur when there are changes in extracellular ionic concentrations (e.g., high potassium) (Gramsbergen et al. 1997) or during a state of reduced cellular energy generation (e.g., hypoglycemia) which reduces the energy-dependent transport systems for kynurenine. Kynurenate formation is reduced also by compounds, such as the metabotropic glutamate receptor (mGluR) agonists and the sulfur-containing endogenous amino acid cysteine sulfinate, which inhibit KAT II (Battaglia et al. 2000; Kocki et al. 2003). Also,  $L-\alpha$ -aminoadipate, a gliotoxic metabolite of the amino acid lysine, decreases the transamination of kynurenine to kynurenate and exerts its neuropathological effects partly by such an action on astrocytes (Wu et al. 1995). In contrast to the effects of the above compounds, pyruvate and other 2-oxoacids increase kynurenate levels by serving as co-substrates for the transamination process (Hodgkins et al. 1999).

Another important neuroprotective metabolite in the kynurenine pathway is picolinic acid, which is produced in neurons from 2-amino-3-carboxymuconic acid semialdehyde. The level of picolinic acid in the CSF declines with age which would contribute to the increasing prevalence of neurodegenerative diseases with age (Coggan et al. 2009). The neuroprotective properties of picolinic acid are achieved in part through its ability to promote chelation of iron and other metals (Testa et al. 1985; Jhamandas et al. 1990; Guillemin et al. 2007).

# 5 Diseases Associated with Kynurenines

#### 5.1 HIV-1-Associated Neurocognitive Disorder (HAND)

Human immunodeficiency virus (HIV-1)-induced neuropathology (ranging from the most severe AIDS–dementia complex to mild cognitive impairment) does not result from direct infection of neurons by the virus but indirectly from the release of neurotoxins from activated immune cells (Glass et al. 1995; Adle-Biassette et al. 1999).

HIV-1 infection causes intestinal mucosal damage and elevated levels of lipopolysaccharides (LPS) in the blood arising from enteric bacteria (Brenchely et al. 2006) Infected monocytes in the blood cross through the blood–brain barrier which is breached in part by the LPS and pro-inflammatory mediators (Zhou et al. 2006). In the CNS, the monocytes not only release viruses that infect other monocytes, but also differentiate into macrophages and release viral proteins, for example, gp120 and Tat that activate microglia and, to a lesser degree, astrocytes. The chemokines and cytokines (e.g., TNF- $\alpha$ , IL-1 $\beta$  and monocyte chemo-attractant protein-1) produced by the activated brain immune cells promote further migration of monocytes across the blood-brain barrier (Boven et al. 2000; Fantuzzi et al. 2003; Persidsky and Gendelman 2003). Additionally, Tat activates IDO in peripheral and brain macrophages - a process that is mediated by p38 MAPK (mitogen-activated protein kinase) and enhanced by IFN- $\gamma$  (Heyes et al. 1992; Sardar and Reynolds 1995; Smith et al. 2001; Fu et al. 2011). The increased expression of IDO eventually results in elevated quinolinic acid levels and neurotoxicity (Fig. 2). Additionally, the released quinolinic acid induces astrocytes to produce more chemokines, thus amplifying the neuropathology (Croitoru-Lamoury et al. 2003). Also, the neuronal damage is exacerbated by increased levels of glutamate due to the latter's impaired reuptake by the damaged astrocytes.

#### 5.2 Alzheimer's Disease

The severity of Alzheimer's disease (AD) depends on the accumulation of two groups of molecules in the CNS – amyloid- $\beta$  fragments and tau proteins. However, there are other pathological processes that contribute to the disorder including excitotoxicity and free radical damage caused by kynurenine pathway metabolites. For example, the kynurenine/tryptophan ratio is increased in AD (Widner et al. 2000), while increased levels of circulating IgA antibodies to IDO have been detected in the blood of AD patients, suggesting an increase in the production of the enzyme (Duleu et al. 2010). Also, amyloid- $\beta$  activates microglia and induces IDO expression, leading to increased quinolinic acid production (Guillemin and Brew 2002; Guillemin et al. 2003a, b) and an associated decreased level of kynurenate (Hartai et al. 2007). NMDAR overactivation by quinolinic acid increases amyloid- $\beta$  production, thus providing a positive feedback for disease progression (Lesne et al. 2005). In rodents, the level of quinolinic acid in the cortex increases with age, providing another mechanism for age-related predisposition to AD.

#### 5.3 Huntington's Disease

Huntington's disease (HD) is an autosomal-dominant disease associated with neuronal loss in the striatum and neocortex. Beal et al. (1986) were the first to report that injection of quinolinic acid to the striatum of rodents produces a pattern of neuronal damage similar to that seen in human postmortem HD tissue. In HD, the evidence points strongly to an over-expression of the abnormal huntingtin gene and consequent neuronal damage by the huntingtin protein (Carter et al. 1999; Usdin et al. 1999). Interestingly, quinolinic acid has also been shown to induce expression of huntingtin in the striatum (Tatter et al. 1995). Additionally, huntingtin increases expression of the NR1/NR2 subunit combination of NMDA receptors in some types of neurons in the striatum, and the resulting increase in cellular sensitivity to quinolinic acid and glutamate (Fan et al. 2007) promotes disease progression.

The huntingtin protein triggers microglial activation (Haass and Selkoe 2007; Thakur et al. 2009) and in the R6/2 mouse, a well-established animal model of HD, the levels of quinolinic acid and 3HK are increased in the cortex and striatum (Guidetti et al. 2006). The increased level of 3HK is due to the combined increased activity of the synthesizing enzyme, KMO, and decreased activity of its degradative enzyme, kynureninase (Giorgini et al. 2005; Sathyasaikumar et al. 2010). Increased 3HK and quinolinic acid levels have also been observed in the striatum and cortex of other animal models of HD, for example, YAC128 and *Hdh*<sup>Q92</sup>/*Hdh*<sup>Q111</sup> mice (Guidetti et al. 2006).

In humans, there is an increased plasma kynurenine:tryptophan ratio in HD patients (Stoy et al. 2005; Forrest et al. 2010). Also, the levels of quinolinic acid and 3HK are increased in the brain of low-grade HD patients, but the levels tend to be similar to those of age-matched controls as the disease progresses (Pearson and Reynolds 1992; Guidetti et al. 2004). The severity of HD is linked to the number of glutamine residues (coded by CAG nucleotide triplets) close to the N-terminal of the huntingtin protein. The increased conversion of tryptophan to kynurenine by IDO in HD patients is further supported by the finding of a positive association between the plasma kynurenine:tryptophan ratio and the severity of HD, as determined by the clinical state or the degree of CAG triplet repeats (Stoy et al. 2005; Forrest et al. 2010). The correlation between kynurenine pathway activity and both disease severity and CAG repeat length represents one of the few known instances of a biochemical parameter which can act as a reliable marker of the disease and, potentially a predictor of disease progression. It also remains a strong possibility that these correlations reflect a causative role for the kynurenine pathway in the severity of the disease (Stone et al. 2012a, b). Thus, one consequence of the extended glutamine repeat sequences in huntingtin may be a change in one or more of the kynurenine pathway enzymes, as a result of which there is an altered susceptibility to neuronal damage.

A more detailed discussion of kynurenines and Huntington's disease, together with some of the issues associated with viewing the kynurenine pathway as a potential therapeutic target for the prevention, delay, and treatment of the disorder, may be found in recent reviews (Stone et al. 2012a, b).

#### 5.4 Parkinson's Disease

Parkinson's disease (PD) is associated with neuro-inflammation: Microglial activation forms part of the pathobiology of PD induced in rodents by 1-methyl-4-phenyl-tetrahydroyridine (MPTP) or LPS (Wu et al. 2002; Liu and Hong 2003). Also, PD patients exhibit microglial activation in the basal ganglia and cortex (Gerhard et al. 2006), with a resulting increase in 3HK and quinolinic acid and subsequent neuronal damage (Ogawa et al. 1992; Zinger et al. 2011). In addition, the antagonist activity of kynurenate at glutamate receptors is reduced in PD, since there is decreased expression of KAT1 in the substantia nigra of a rodent model of PD (Knyihar-Csillik et al. 2004) and there are lowered levels of kynurenate in the basal ganglia and cortex of PD patients (Ogawa et al. 1992). Overall, kynurenine pathway activation in PD seems to be such that the production of neurotoxic metabolites (quinolinic acid, 3HK) by microglia and infiltrating microphages outweighs the production of neuroprotective metabolites kynurenate (by astrocytes) and picolinic acid (by neurons) (Zinger et al. 2011) (Fig. 2).

# 5.5 Schizophrenia

Patients with schizophrenia have increased plasma levels of the pro-inflammatory cytokines IFN- $\gamma$ , TNF- $\alpha$ , and IL-6 (Kim et al. 2009) and an associated activation of the kynurenine pathway (Barry et al. 2009). There is increased expression of IDO which is accompanied by a relatively greater activation of astrocytes than microglia, thus leading to the observed increase in levels of kynurenate in the cortex and CSF (Schwarcz et al. 2001; Erhardt et al. 2001; Muller et al. 2011). The blockage of NMDARs by kynurenate in the cortex could account for the glutamatergic hypofunction that is now believed to be a major factor in creating the established dopaminergic dysfunction in schizophrenia. For example, since glutamate and nicotinic acetylcholine receptors (nAChRs) are essential for cognitive processing, their blockage by kynurenate explains the cognitive deficit in schizophrenis (Zmarowski et al. 2009; Wu et al. 2010). Additionally, the sensory gating deficits in schizophrenia can be explained by kynurenate decreasing excitability of CA1 stratum radiatum interneurons via its blockade of NMDARs and nAChRs (Alkondon et al. 2011).

#### 5.6 Depression

Depression is associated with an activated immune system (Maes et al. 1995; Maes 1999). For example, pro-inflammatory cytokines produced by activated macrophages induce symptoms of depression when given to volunteers (Smith 1991; Dantzer et al. 2011) and patients receiving IFN- $\alpha$  for hepatitis C infection develop symptoms of depression. Also, depression is associated with increased IDO activity (which is reflected in an increased plasma kynurenine:tryptophan ratio), and a reduced level of the neuroprotective compound kynurenate (Wichers et al. 2005; Myint et al. 2007). The activity of TDO also is increased by pro-inflammatory cytokines, and by cortisol (Leonard and Myint 2006). Hence, the increased levels of cortisol during depression augment the activation of the kynurenine pathway by the cytokines to cause the neurodegenerative changes in susceptible brain regions – such as the subgenual anterior cingulate cortex, hippocampus, and amygdala – that are linked to the symptoms of depression.

Depression has an established association with low levels of serotonin for synaptic transmission, a link which has led to the widespread use of serotonin uptake inhibitors for treating depression. However, the diversion of tryptophan to the kynurenine pathway following immune system activation is likely to reduce the availability of tryptophan for serotonin synthesis. Changes of serotonergic neurotransmission may, therefore, be secondary consequences of kynurenine pathway activity, a hypothesis that could partly explain the association between activation of the kynurenine pathway and depression (Mackay et al. 2009).

#### 5.7 Stroke

Cerebral ischemia induced by 10 min of bilateral carotid occlusion in gerbils results in increased levels of brain quinolinic acid at 4 days post ischemia (Heyes and Morrison 1997). In stroke, there is increased production of pro-inflammatory cytokines including IFN- $\gamma$ , IL-1 $\beta$ , IL-8, IL-6, and IL-17 (Tarkowski et al. 1995; Kostulas et al. 1999; Hoshi et al. 2009); a subsequent localized increase in IDO expression (Taguchi et al. 2008; Hoshi et al. 2009); and increased kynurenine metabolites in the brain (Saito et al. 1993).

In humans, activation of the kynurenine pathway in stroke is supported by the finding of an association between stroke and an early onset decrease in plasma tryptophan levels and 3-hydroxyanthranilic acid:anthranilic acid ratio. Stroke also causes increases in the plasma kynurenine:tryptophan ratio (a measure of IDO expression), neopterin (a major inflammation marker), S100B protein (a marker of brain injury), and lipid peroxidation products (malondialdehyde and 4-hydroxynonenal ). The levels of S100 and peroxidation products, as well as aspects of kynurenine metabolism, correlate with the infarct volume, as determined by CT scans (Darlington et al. 2007). Furthermore, the degree of IDO expression is well correlated with mortality risk following a stroke (Darlington et al. 2007) and with the subsequent degree of cognitive impairment following acute stroke (Gold et al. 2011). The latter observation is consistent with data showing very strong correlations between kynurenine pathway activity and cognitive function following cardiac bypass or thoracic surgery, to the extent that the biochemical changes may be used to predict cognitive outcome (Forrest et al. 2011).

#### 5.8 Multiple Sclerosis

In multiple sclerosis (MS), there are parenchymal infiltrates associated with the multifocal areas of demyelination, macrophages and microglia being the main cell types found in the infiltrates. The activated macrophages and microglia release cytokines (IL-6, TNF- $\alpha$ , IL-1 $\beta$ , IL-12 and IL-23) that are capable of activating the kynurenine pathway.

Patients with MS have low tryptophan levels in serum and CSF (Sandyk 1996; Rudzite et al. 1996), suggesting activation of the kynurenine pathway. There have also been reports of decreased kynurenate levels in the CSF (Rejdak et al. 2002) and increased levels of plasma kynurenate (Hartai et al. 2005). The difference is unlikely to be due to the sample used since kynurenate easily crosses the blood– brain barrier. However, the difference might be due to the stage of the disease with increased kynurenate production during the active phase and a decreased production during the chronic non-active phase (Lim et al. 2010).

It seems that activation of microglia and perivascular macrophages in MS causes increased amounts of quinolinic acid that overwhelm the neuroprotective actions of kynurenate and contribute to the axonal demyelination. The oligodendrocytes that are damaged in MS lack IDO-1 activity and hence play an insignificant role in the kynurenine pathway; however, they do have NMDAR on their processes and AMPA/kainate receptors on the cell bodies (McDonald et al. 1998), making them susceptible to quinolinic acid– and glutamate-mediated excitotoxicity.

#### 6 Therapeutic Possibilities

Acyclovir, an established drug against herpes virus infection, has been shown to inhibit quinolinic acid–induced lipid peroxidation in rat brain (Muller et al. 2005a), making it a potential treatment for quinolinic acid–mediated neurodegenerative diseases even when viral infection is not a primary factor.

A number of other compounds such as Necorostatin-1, Annulin, and Exigamine A have been shown to be effective IDO inhibitors in experimental models (Muller et al. 2005b; Brastianos et al. 2006; Pereira et al. 2006). That activity may make the compounds suitable for use as anticancer therapy or to inhibit the production of quinolinic acid, 3HK, and 3-hydroxyanthranilic acid in order to reduce excitotoxicity. However, they have not yet been shown to effective and safe enough to enter general clinical use (Takikawa 2007).

Because kynurenate production takes place in astrocytes, whereas 3HK and quinolinic acid are produced in microglia, the two branches of the kynurenine pathway are anatomically segregated. Additionally, these two branches are functionally segregated since the enzymes KAT II and KMO can be blocked selectively: Blocking of KAT II has no effect on 3HK/quinolinic acid levels, while blocking KMO decreases 3HK/quinolinic acid levels but increases kynurenate levels (Amori et al. 2009). Selective KMO inhibitors have been shown to produce neuroprotection against neuronal cell death in hippocampal slices (Carpenedo et al. 2002), brain

ischemia in gerbils (Cozzi et al. 1999), and dystonia in hamsters (Hamann et al. 2008). The therapeutic use of KMO inhibitors as neuroprotectants may be particularly effective when immune system activity is responsible for the changes in kynurenine pathway activation. Examples of this situation include parasitic disorders with CNS involvement. Infection of appropriate strains of mice with *Plasmodium* species, able to produce cerebral involvement similar to that encountered in humans, leads to ataxia and death within a few days; KMO inhibition reduces symptom severity and prevents mortality (Clark et al. 2005). Protection against brain damage has also been shown in a murine model of cerebral trypanosomiasis ("sleeping sickness"), a disease which is almost invariably fatal in humans (Rodgers et al. 2009). Despite these successes, the compounds used to date as KMO inhibitors penetrate the blood–brain barrier only poorly, and even greater success is likely to be achieved with the development of compounds that effectively cross this barrier more effectively (Zwilling et al. 2011).

# 6.1 Diet

Finally, it should be noted that in the fields of neurodegeneration and neuroprotection, as with several other major areas of central and peripheral pathology, dietary and nutritional factors may not only play a part in determining the extent of brain damage, but may also point toward potential therapeutic strategies for the reduction of damage. For example, there is accumulating evidence that a ketogenic diet (high in fat, low in carbohydrate) can increase plasma levels of acetoacetate,  $\beta$ -hydroxybutyrate, and acetone. The high levels of these ketone bodies may then provide neuroprotection against a number of CNS disorders in which brain damage occurs (in animal models or patients) including epilepsy, Alzheimer's disease, Parkinson's diseases, stroke, and traumatic brain injury (Gasior et al. 2006). Mice on a 4-week ketogenic diet exhibited smaller increases in brain TNF- $\alpha$  induced by the potent exogenous excitotoxin kainate (Jeong et al. 2011); the dietary effect would lead to reduced production of endogenous excitotoxins. Other studies have reported that ketone bodies induce the expression of KAT I and II in glial cells (Chmiel-Perzynska et al. 2011), leading to increased kynurenate levels in areas such as the hippocampus and striatum (Zmarowski et al. 2009).

Since some lifestyle activities including fasting or strenuous exercise can mimic ketogenic diets by promoting the formation of ketone bodies, it is likely that these or similar interventions will be examined closely as potential means by which the incidence and severity of CNS damage can be minimized, or recovery enhanced.

# 7 Conclusion

The key message from the work described above is that pharmacological interference with the kynurenine pathway should have importance in the treatment of a wide range of neuro-inflammatory and neurodegenerative disorders, including those with primarily neurological or psychiatric symptomatology. This strategy would represent a major conceptual advance over the use of general receptor antagonists, since glutamate is the most widely used excitatory neurotransmitter in the mammalian CNS. A general blocker of glutamate receptors (or the major subdivisions of NMDA, AMPA, and metabotropic receptors) is, as a result, likely to induce a plethora of unwanted side effects that may include neurological manifestations such as paralysis, loss of memory, and compromised function of controlling output systems to the cardiovascular and respiratory systems. Interference with the kynurenine pathway should only affect those neurons and glia affected by the brain insult and thus should produce a locally directed modification of dysfunction with correspondingly fewer associated problems – one of the gold standard objectives of modern drug development.

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# **Role of Neurotoxicity in Depression**

Irena Smaga, Weronika Krzyżanowska, Bartosz Pomierny, Bogusława Budziszewska, Andrzej Pilc, and Gabriel Nowak

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I. Smaga • W. Krzyżanowska • B. Pomierny

Department of Toxicology, Jagiellonian University Medical College, Kraków, Poland

B. Budziszewska

Department of Toxicology, Jagiellonian University Medical College, Kraków, Poland

Institute of Pharmacology, Polish Academy of Sciences and Center of Excellence in Neuropsychopharmacology, Kraków, Poland

A. Pilc

Institute of Pharmacology, Polish Academy of Sciences and Center of Excellence in Neuropsychopharmacology, Kraków, Poland

Jagiellonian University Medical College, Kraków, Poland

G. Nowak (🖂)

Department of Pharmacobiology, Jagiellonian University Medical College, Kraków, Poland

Institute of Pharmacology, Polish Academy of Sciences and Center of Excellence in Neuropsychopharmacology, Kraków, Poland e-mail: nowak@if-pan.krakow.pl

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#### Abstract

Despite years of research, the pathophysiology of depression and the mechanism of action in antidepressant drugs largely remain unknown. The subsequent hypotheses examining depression witnessed attention being paid to the decrease of the serotonin and noradrenaline synaptic concentrations, adaptive changes in the neurotransmitters receptor, activation of the immune system, hyperactivity of the hypothalamic-pituitary-adrenal (HPA) axis, and enhanced glutamatergic activity. Current research suggests that disturbed neuronal plasticity and the degeneration of neurons and glia cells may be involved in the pathogenesis of depression. In fact, in depressed patients, structural changes are observed in several brain regions, mainly in the hippocampus. Also, in various animal models of depression, morphological alteration in neurons and glial cells is present besides depression-like behavioral changes. A lot of evidence indicates that an increased amount or activity of glucocorticoids, glutamate, and proinflammatory cytokines is the reason for these changes. In both animal models of depression and postmortem studies, a reduced neurogenesis in the dentate gyrus of the hippocampus; a decrease in the total number of neurons and astrocytes; a reduction of the dendritic length, branching density, or the number of synapses; and a decrease in the brain-derived neurotrophic factor were observed. Moreover, activation of microglia, a source of proinflammatory cytokines and reactive oxygen species, has a disadvantageous effect on neurons and astrocytes. However, it should be noted that not all of the studies demonstrate neurodegenerative changes in depression, or the glucocorticoids' detrimental action, so further studies are needed in order to clearly determine the contribution of neurotoxic agents in the pathogenesis of depression and also the importance of the neuroprotective properties of some antidepressant drugs in their therapeutic effect.

#### Keywords

Animal models of depression • Antidepressant • Depression • Neurotoxicity

5HTSerotoninACTHAdrenocorticotropic hormoneBDNFBrain-derived neurotrophic factorCMSChronic mild stressCREBCyclic AMP response element-binding proteinCRFCorticotropin-releasing factor	List of Abbreviations		
BDNFBrain-derived neurotrophic factorCMSChronic mild stressCREBCyclic AMP response element-binding protein	5HT	Serotonin	
CMSChronic mild stressCREBCyclic AMP response element-binding protein	ACTH	Adrenocorticotr	ropic hormone
CREB Cyclic AMP response element-binding protein	BDNF	Brain-derived n	eurotrophic factor
	CMS	Chronic mild st	ress
CRF Corticotropin-releasing factor	CREB	Cyclic AMP res	sponse element-binding protein
	CRF	Corticotropin-re	eleasing factor

C-reactive protein
Chronic unpredictable stress
Glial fibrillary acidic protein
Hypothalamic-pituitary-adrenal
Lipopolysaccharide
Medial prefrontal cortex
Noradrenaline
<i>N</i> -methyl-D-aspartate
Nitric oxide synthase
Prefrontal cortex
Reactive oxygen species
Selective serotonin reuptake inhibitors
Tricyclic antidepressant

#### 1 Introduction

The first effective antidepressant drugs were introduced around 50 years ago, although their mode of action remained unclear (Charney et al. 1981). The investigation into the mechanism of action of antidepressants (iproniazid, imipramine) showed that these substances increase the central serotonin and norepinephrine transmission. Using these results as the basis, it was hypothesized that the cause of depression is the decrease in the level of monoamines such as serotonin or norepinephrine in the synaptic space, as well as a reduction in nerve conduction (Berton and Nestler 2006). It has been also shown that antihypertensive drugs such as reserpine, which cause monoamine depletion, sometimes lead to depression (Freis 1954). However, the hypothesis concerning monoamine has been recently questioned. It has been observed that not all drugs that increase monoamine concentration exert an antidepressant effect, and drugs such as monoamine oxidase inhibitors (tranylcypromine), and inhibitors of selective serotonin reuptake (fluoxetine), act immediately through the enhancement of monoaminergic conduction, yet their therapeutic effect in the improvement of mood is observed after a few weeks of therapy (Ruhé et al. 2007). Both the clinical and etiological heterogeneities of depression make it hard to define its pathophysiology (Hasler 2010). The last few years have paid witness to a variety of theories explaining the etiology of this disease. A decrease in the synaptic levels of monoamines, changes in neurotransmitter receptors, protein kinases, proinflammatory cytokines, neurotrophic factors, and hyperactivity of the hypothalamic-pituitary-adrenal (HPA) axis in the pathophysiology of depression were postulated as being potentially involved. Studies conducted in recent years have provided abundant evidence that the disturbance of molecular and cellular regulation of neuronal plasticity may be involved in the pathogenesis of depressive disorders (Nestler et al. 2002; Pittenger and Duman 2008). A considerable amount of data has suggested that degeneration of neurons and glia cells evoked by a disturbance in neurotrophic factor pathways and enhanced glutamate, glucocorticoids, and proinflammatory cytokines' action play an important role in the pathogenesis of this disease. However, it has not presently been conclusively established whether morphological and functional brain changes observed in depression are one of the causes of disease or if they are the result of depression.

# 2 The Evidence for Brain Structures Changes in Depression: Clinical Data

Several changes have been noted in the structures anatomically linked with depression, such as the prefrontal cortex, hippocampus, amygdala, or nucleus accumbens. First of all, the structure which is the most vulnerable to atrophy is the hippocampus. The hippocampus is an area of the brain responsible for learning, memory, mood, and emotions. It plays an important role in the consolidation of information from short-term memory to long-term memory and spatial orientation. Volume reduction of this structure in depression, observed in postmortem studies, might be the result of many processes, including the remodeling of cellular elements, retraction of dendrites, reduction of neurogenesis in the dentate gyrus, and loss of glial cells (McEwen 1999; Rajkowska 2000; Czéh et al. 2001). In depressed patients, it was observed that the size of pyramidal neurons decreased, whereas the packing density of glial cells, pyramidal neurons, and granule cell neurons was significantly greater (Stockmeier et al. 2004; Tsopelas et al. 2011). The hippocampus is the most extensively studied region in depression by neuroimaging, and the resulting findings seem to suggest that hippocampal volumetric reductions are associated with depression (Bremner et al. 2000; Frodl et al. 2002b; MacQueen et al. 2003).

There were also reports that the volume of other brain structures have also decreased in postmortem research. Patients with depression had a significantly smaller medial orbitofrontal cortical volume, without smaller volumes of other frontal regions. These findings are consistent with other studies where the reduced volume of the orbitofrontal cortex, as well as the reduction of the entire frontal lobe, was shown (Bremner et al. 2002; Shah et al. 2002; Frodl et al. 2006). This change occurs in a more severely ill patient group, but not in those who were less severely ill (Bremner et al. 2000; Lacerda et al. 2004). Similar to the hippocampus, the decreased volume of the orbitofrontal cortex is caused by an increased density of glial cells and neurons (Rajkowska 2000). Depression was also associated with neuronal loss in some of the subcortical structures investigated (nucleus basalis, substantia nigra, raphe nucleus) (Tsopelas et al. 2011).

Examination of the brain of depressives showed an enlargement of the cerebral ventricles (Kellner et al. 1986). With this increase in ventricular volume, it was noted that the surrounding tissue shrinks (mammillary bodies, thalamus) and there are changes in formations that are connected to structures that pass through the third ventricle, such as the hippocampus (via the fornix) and the amygdala, which is connected via the stria terminals (Baumann et al. 1997; Dasari et al. 1999; Bernstein et al. 2008).

There were also noted changes in the volume of the amygdala. The amygdala is a structure that contains an extensive system of neuronal connections. It is involved in the analysis and processing of incoming information and with then comparing them to the stored patterns of emotional memory. Patients with depression have impaired functioning in emotional tasks involving the amygdala and, furthermore, they have an abnormal resting amygdala blood flow (Ledoux 2000; Phan et al. 2002). Depressed patients had reduced amygdala core nuclei volumes, without any differences in the total amygdala volumes. Studies indicate that reduction of the hippocampal volume was often accompanied by the volume reduction of amygdala (Caetano et al. 2004). It seems that glucocorticoid-induced neurotoxicity has been mediated by the glutamatergic connections in these closely cooperating structures (Sheline et al. 1998). It was noted that the initial stage of the disease was characterized by increased volume of this structure (Frodl et al. 2002a, b; Lange and Irle 2004; Weniger et al. 2006), and during the development of the depression, the volume reduction was observed, especially in females (Bremner et al. 2000; Caetano et al. 2004; Hastings et al. 2004).

# 3 The Evidence for Changes in Brain Structures in Depression: Experimental Data

Depression is a very diverse disease, with mostly different symptoms amongst individuals, and its etiology is still poorly recognized despite extensive investigation (Table 1). More importantly, it is not possible to reproduce some of the symptoms characteristic of major depression in animals, such as suicidal thoughts, a depressed mood, or the feeling of guilt. This is the reason why it is hard to delineate the most accurate animal model of depression.

The majority of the presently applied models are based on different stress paradigms and in different stress patterns (chronic, acute, unpredictable). Chronic mild stress (CMS), which consists of the long-term exposure of animals to a series of unpredictable, mild stressors, produces depression-like behavioral changes, including anhedonia. In this model of depression, a decrease in the total number of granule cells in the ventral hippocampus, an enhanced blood corticosterone level, and an increase in the number of caspase-3 positive neurons in the cerebral cortex were observed (Jayatissa et al. 2006; Bachis et al. 2008; Nowak et al. 2010). In chronic unpredictable stress (CUS), which has been conducted on rabbits, glial atrophy in the PFC has been confirmed (Banasr and Duman 2008). In prenatal restraint stress, which, similar to CMS, exhibits face, predictive and construct validity, a decrease in hippocampal cell neurogenesis in the dentate gyrus and impairment in the hippocampal-related spatial tasks were indicated (Lemaire et al. 2000). A decrease in hippocampal neurogenesis has also been observed in the olfactory bulbectomy (Jaako-Movits et al. 2006). In another examination, with chronic or acute restraint stress, there were a decrease in the total number of neurons and the structure volume and a decrease of the neurogenesis of the CA3 field dentate gyrus (Pham et al. 2003). In chronic, social stress in tree shrews,

Animal model of depression	Structural changes	References	
Chronic mild	▼Granular cell number in hippocampus	(Bachis et al. 2008; Nowal	
stress	▲Caspase cascade in cerebral cortex cells et al. 2010)		
Chronic	▲Atrophy of the glial cells in prefrontal cortex	(Banasr and Duman 2008; Nowak et al. 2010)	
unpredictable stress	▼Granular cell number in hippocampus		
Tree shrews	▼Hippocampal volume	(Czéh et al. 2001; Lucasser et al. 2001)	
	▼Neurogenesis in hippocampus		
	▲Apoptosis in the temporal cortex		
Restraint stress	▼Neurogenesis in the region CA3 of the hippocampus	(Pham et al. 2003)	
	▼Total neuron number in the hippocampus		
	▼Hippocampal volume		
Maternal	▲Apoptosis in the hippocampus	(Lee et al. 2001)	
separation	▼Cell proliferation in the hippocampus		
Learned helplessness	▼Cell proliferation in dentate gyrus	(Vollmayr et al. 2003)	
Olfactory bulbectomy	▼Hippocampal neurogenesis	(Jaako-Movits et al. 2006)	
Prenatal stress	▼Hippocampal cell neurogenesis	(Lemaire et al. 2000; Murmu et al. 2006)	
	▼Dendritic length and complexity in prefrontal cortex		

 Table 1
 Summary of structural alterations related to neurodegenerative processes demonstrated in animal models of depression

a model with high validity for depression, a reduced hippocampal volume, a decreased hippocampal neurogenesis, and an increased number of apoptotic cells in the temporal cortex occur (Czéh et al. 2001; Lucassen et al. 2001). A decrease in cell proliferation and enhanced apoptosis in the dentate gyrus of the hippocampus were reported in the maternal separation rat model (Lee et al. 2001). The expression of antiapoptotic proteins (Bcl-2, Bcl-xL) was lower in the cingulate and frontal cortex, hippocampus, and central nucleus of the amygdala in CUS (an animal model of depression) (Kosten et al. 2008). In the restraint model of psychological stress in mice, the increased proliferation of microglia cells was observed (Nair and Bonneau 2006).

# 4 The Involvement of the HPA Axis in Neurodegenerative Processes in Depression

Amongst the possible mechanisms involved in the brain's morphological changes observed in depression, hyperactivity of the HPA axis, itself leading to prolonged hypercortisolemia, has been considered. This mechanism may play an important role in the volumetric and neurodegenerative alteration of several brain areas, especially in those with a high concentration of glucocorticoid receptors, such as the hippocampus (Herman et al. 2003). It has been observed that, on the one hand, stress-induced neurotoxicity can cause volumetric changes in different brain structures, while, on the other hand, these changes may in turn induce an increase in HPA axis activity (Cerqueira et al. 2005). This results from the fact that stress and glucocorticoids induce neuronal loss mainly in the hippocampus, a structure which exerts an inhibitory effect on HPA axis activity. Thus, hippocampal damage disinhibits the HPA axis and, as a result, additionally enhances glucocorticoid synthesis and release. This "glucocorticoid cascade hypothesis," formulated by Sapolsky et al. (1986), was proposed as an explanation of hippocampal damage in both some neurodegenerative disorders and depression.

Psychological studies conducted in patients affected by depression at the end of the 1970s revealed the role of psychological stress as a factor contributing to the development of depression (Brown and Harris 1978). At the same time, HPA axis abnormalities, including an increased cortisol serum level, were shown to be one of depression's features. In addition, it has been observed that dexamethasone administration suppresses the cortisol level in the blood of depressed patients less potently than in healthy people (Carroll 1982). The participation of the HPA axis dysfunction in the pathogenesis of depression has been also supported by the observations that patients suffering from Cushing's syndrome, characterized by elevated levels of serum cortisol, often exhibit symptoms of depression (Sonino et al. 1993). Similarly, it has been shown that the administration of chronic glucocorticoids evoked depression-like symptoms in rodents (Gourley et al. 2007). A HPA axis dysfunction in depression can manifest in a variety of ways, including excessive sensitivity of the adrenal gland to the adrenocorticotropic hormone (ACTH) (Parker et al. 2003), increased secretion of the corticotropin-releasing factor (CRF) (Nemeroff and Owens 2002), or the reduction in negative feedback exerted via the glucocorticoids' receptor (Brown et al. 2004).

Glucocorticoids regulate metabolism and immune system activity and are essential for maintenance, homeostasis, and adaptation to stress, but their long-term increased level or disturbance in the circadian rhythm can have a deleterious effect on the brain's morphology and function. Glucocorticoid receptors are present in many brain regions which are affected by depression, such as the prefrontal cortex (PFC), hippocampus, nucleus accumbens, striatum, limbic system, and ventral midbrain (Herman 1993; Van Craenenbroeck et al. 2005). It was discovered that when corticosterone concentration increases for a long time, it causes neuronal damage in the CA1 and CA3 hippocampal region and enhances neurodegenerative processes evoked by excitatory amino acids, hypoxia, hypoglycemia, seizures, or age. One of the long established changes caused by chronic stress or long-term administration of glucocorticoids is the reorganization of the apical dendrites of pyramidal cells in the CA3 region of the hippocampus. Stress or glucocorticoid treatment induces an extensive reduction of the total dendritic length, a reduction in the distal dendritic branching density, and a reduction of the number of synapses in the hippocampal CA3 region in rats (Magariños et al. 1999; Sousa et al. 2000). Apart from the hippocampus, atrophy of the dendrites was also witnessed in the prefrontal cortex in experimental animals after chronic stress (Magariños and McEwen 1995; Wellman 2001; Duman 2002).

It seems that one of the reasons for hippocampal volume reduction observed in depression is an impairment in the process of neurogenesis. A lot of data indicated that glucocorticoids, stress, and glutamate inhibited neurogenesis in the dentate gyrus of the hippocampus, while physical and mental activity, insulin-like growth factor, serotonin, estradiol, and adrenalectomy enhanced this process in experimental animals. It seems that glucocorticoids and glutamate regulate neurogenesis by a common mechanism, because *N*-methyl-D-aspartate (NMDA) receptor antagonists inhibit corticosterone action, while NMDA activation blocks the effect of adrenalectomy (Cameron et al. 1998). Since glucocorticoids enhance extracellular glutamate concentration in the hippocampus, their inhibitory effect on the process of neurogenesis may result from the enhancement of glutamate activities.

The mechanisms of glucocorticoids' deleterious action on brain cells are unclear, but most probably are connected with enhancing glutamate action, inhibition of astrocyte glucose uptake, and decreasing the synthesis of neurotrophic factors. For example, stress or glucocorticoids reduce synthesis of the brain-derived neurotrophic factor (BDNF), a compound responsible for the growth, maturation and survival of nerve cells as well as for the processes of arborization and brain plasticity. It has been shown that chronic stress decreases the synthesis of BDNF in the hippocampus, whereas antidepressants lead to an increase in its production in the hippocampus and PFC (Shimizu et al. 2003). It has also been found that stress reduces the BDNF and tyrosine kinase B (TrkB) (BDNF-specific receptor) genes' expression and phosphorylation of the cyclic AMP response element-binding protein (CREB), which is the transcription factor that regulates the expression of BDNF (Nibuya et al. 1995, 1996; Duman et al. 1997, 2000).

The role of glucocorticoids in the pathogenesis of depression is evidenced by the data indicating the antidepressant effect of GR antagonists and glucocorticoid synthesis inhibitors in animal models (Baez and Volosin 1994; Healy et al. 1999). Clinical data also showed the antidepressant action of glucocorticoid synthesis inhibitors, and clinical trials with specific glucocorticoid and CRF receptor antagonists as potential antidepressants are in progress (Wolkowitz and Reus 1999; Mathew et al. 2008).

# 5 The Involvement of Enhanced Glutamate Transmission in Neurodegenerative Processes in Depression

The hypothesis that dysfunction of the glutamatergic system may be involved in the pathophysiology and treatment of depression was proposed in 1996 (Skolnick et al. 1996), following several discoveries concerning the importance of the NMDA receptor in antidepressant action (the first article by Trullas and Skolnick evidenced that NMDA receptor functional antagonists exhibit antidepressant-like activity, Trullas and Skolnick 1990). Many studies demonstrated that patients with depression have an elevated glutamate level in the blood (Kim et al. 1982; Altamura et al. 1993) and a positive correlation between the blood glutamate level and the severity of depressive symptoms was discovered (Mauri et al. 1998; Mitani et al. 2006). Similarly, it was

demonstrated that the glutamate level in the cerebrospinal fluid of patients with depression is higher than in the healthy controls (Levine et al. 2000). The disturbance of glutamatergic neurotransmission in depression is also supported by postmortem studies. Alterations in [<sup>3</sup>H]CGP39653 (an antagonist of the glutamate binding site of the NMDA receptor) binding and its modulation by glycine were demonstrated in both the frontal cortex and hippocampus in suicide/depressed humans (Nowak et al. 1995; Beneyto et al. 2007). Furthermore, a decrease of [<sup>3</sup>H]L-689,560 binding (an antagonist of the NMDA receptor glycine site), as well as the reduction in the NR1 subunit of NMDA receptor immunoreactivity in the superior temporal cortex of depressed patients was reported (Nudmanud-Thanoi and Reynolds 2004). Also, changes in mRNA/protein concentrations of the glutamate receptor subunits (NMDA, metabotropic) in the brain regions of depressed subjects were described (e.g., Law and Deakin 2001; Feyissa et al. 2009; Deschwanden et al. 2011).

Elevated concentration of extracellular glutamate is considered to be the main reason for the degeneration of neurons and/or cell death observed in neurodegenerative diseases (Hardingham and Bading 2010). Environmental stress, which increases both glutamate and glucocorticoids, has a huge impact on the morphology of the brain tissue and is also a risk factor for mood/anxiety disorders (de Kloet et al. 2005). Studies in rats using the microdialysis technique indicated an increase in extracellular levels of excitatory neurotransmitters (glutamate and aspartate) within the striatum, hippocampus, nucleus accumbens, and medial prefrontal cortex (mPFC) after acute restraint stress (for 20 min). The highest glutamate level increase was observed in the mPFC, when compared to other structures. The same result was obtained in swimming stress, conducted in the same period of time. Glucocorticoids, which inhibit glutamate uptake into glia cells, are probably responsible for an enhanced intracellular glutamate level and, consequently, the increase of excitotoxicity processes. It has been reported that an elevated level of glucocorticoids in the hippocampus and mPFC increased glutamate concentration and that both of these compounds are responsible for the retraction of dendrites in the CA3 region of the hippocampus (McEwen 2005; Pittenger and Duman 2008; Gorman and Docherty 2010). In glutamatergic synapses, the typical mechanisms of the synaptic plasticity affecting their activity are changes in the shape and size of the synaptic shafts and spines (Sorra and Harris 2000). Stress can affect the processes of the synaptic plasticity of various brain structures, and different forms of stress may induce atrophy, retraction, and remodeling of dendrites of the pyramidal neurons in the hippocampal CA3 region (Watanabe et al. 1992; Magariños and McEwen 1995). It has been demonstrated that certain intracellular processes, associated with glutamatergic receptor stimulation, may be involved in the pathophysiology of depression and neuronal degeneration. Stimulation of the NMDA receptor leads to an increase in intracellular calcium concentration in postsynaptic neurons. Ca<sup>2+</sup> bind to and stimulate the Ca<sup>2+</sup>/calmodulin-dependent protein kinase, which via action on the nitric oxide synthase (NOS) may then enhance the synthesis of the nitric oxide (II) (Southam and Garthwaite 1993; Li and Poulos 2005). Nitric oxide is a membrane-permeable molecule associated with the processes of cellular signaling and communication in various systems. In the brain, nitric oxide regulates many physiological processes, including neurotransmission, synaptic plasticity, neuromorphogenesis, gene expression, and the perception of pain and learning (Esplugues 2002), suggesting that its enhanced level may contribute to the etiology of depression. Clinical studies have shown increased immunoreactivity for nNOS in the hippocampal CA1 and subiculum region in patients suffering from major depression and bipolar disorder (Oliveira et al. 2008). Administration of 7-nitroindazole, an inhibitor of neuronal nitric oxide synthase (nNOS), into the dorsal hippocampus, induces an antidepressant-like effect in rats (Joca and Guimarăes 2006). Moreover, paroxetine, a selective serotonin reuptake inhibitor, inhibits the activity of constitutive nitric oxide synthase (cNOS) (Finkel 1996).

# 6 The Involvement of the Immune System in Neurodegenerative Processes in Depression

Patients suffering from depression have a higher incidence of immune system disorders when compared to the total population, and also depression often occurs as a side effect of cytokine therapies that are used in the treatment of hepatitis C, multiple sclerosis, Kaposi's sarcoma, and many other cancer types. Depression occurs most frequently in the rapies using IFN- $\alpha$  and IL-2, less frequently during IFN- $\beta$  administration, and has not been recorded in the case of IFN- $\gamma$  (Valentine et al. 1998; Gohier et al. 2003). Patients with depression often exhibit increased activity of the immune system, which is reflected in the increase of the level of "positive" acute phase proteins - C-reactive protein (CRP), haptoglobin, ceruloplasmin - as well as in the reduction of the "negative" acute phase proteins' concentration (albumin, transferrin, retinol-binding protein). Depression is also more common in patients suffering from disease connected with immune component disturbances (Kronfol 2002). Studies conducted in experimental animals have shown that administration of lipopolysaccharide (LPS) or IL-1 evoked "sickness behavior" syndrome, which has symptoms that are similar to those observed in depression (Kent et al. 1994). Similarly, administration of LPS to healthy volunteers resulted in the appearance of symptoms of anxiety and depression (Reichenberg et al. 2001). A lot of data especially indicates the important role of TNF- $\alpha$ , IL-1, and IL-6 in the pathogenesis of affective disorders. Under physiological conditions, these cytokines positively affect the central nervous system, through neurogenesis enhancement and trophic action on neurons. However, excessive stimulation of signal transduction pathways through the proinflammatory cytokines in the central nervous system can have a disadvantageous effect (Miller et al. 2009). Peripheral administration of LPS causes an increase in the concentration of TNF- $\alpha$  and IL-1 in the hippocampus, which is associated with a decreased expression of BDNF and TrkB in this brain structure. This leads to a decrease in the neurogenesis processes, impairs cognitive function and may also cause an increase in glutamatergic transmission, enhances oxidative stress, and induces apoptosis of astrocytes and oligodendrocytes, thus, the processes that are most likely involved in the pathogenesis of depression (Wu et al. 2007; Miller et al. 2009). Apart from enhancing neurodegenerative processes, proinflammatory cytokines can modify the level of neurotransmitters. Animal studies suggest that proinflammatory cytokines alter the turnover of serotonin (5HT) in different brain regions (Dunn et al. 1999), lead to the downregulation of serotonin receptor  $5HT_{1A}$ (Cai et al. 2005), and inhibit serotonin synthesis through activation of the indoleamine 2,3-dioxygenase enzyme that converts the tryptophan (precursor of serotonin) into kynurenine (Schwarcz and Pellicciari 2002). Additionally, proinflammatory cytokines, especially IFN- $\alpha$ , increase the kynurenine-3monooxygenase enzyme and in this way shift kynurenine metabolism toward the synthesis of 3-hydroxykynurenin and quinolinic acid, NMDA receptor agonists which can contribute to the neurotoxic processes observed in depression (Zunszain et al. 2012). Cytokines also affect dopamine neurotransmission by upregulation of the dopamine receptor and an increase of the nitric oxide (II), which in turn inhibits the synthesis of dopamine (Kitagami et al. 2003; Moron et al. 2003). Proinflammatory cytokines were also indicated as modulating the function of the HPA axis. For example, administration of IL-1 and IL-6 enhances CRF secretion and consequently activates the HPA axis (Besedovsky and del Ray 1996). Cytokines can activate the HPA axis acting on the hypothalamus, pituitary, or adrenal cortex and can also modulate glucocorticoid receptor action (Pariante et al. 1999).

# 7 The Disturbed Glucose Metabolism in Depression

Functional neuroimaging research of depressed patients showed anatomical differences in the glucose metabolism in the brain. In 1985, and for the first time, scientists described reduced glucose metabolism in the prefrontal cortex in bipolar depressives (Baxter et al. 1985). These findings were confirmed in other investigations, which then also demonstrated hypometabolism in unipolar patients (Baxter et al. 1989; Hurwitz et al. 1990; Martinot et al. 1990). Further reports indicate the coexistence of reduced blood flow and impaired glucose metabolism in the prefrontal cortex in patients with a reduced cortical volume (Drevets et al. 1997). Moreover, alterations of brain glucose metabolism are stronger in patients with hyperactivity of the HPA axis, which suggests that the impaired glucose metabolism can result from elevated glucocorticoid action.

On the other hand, numerous studies have shown a correlation between an increased severity of depressive symptoms and increased use of glucose in the subgenual anterior cingulate cortex (Drevets et al. 1999; Osuch et al. 2000), although the opposite result has also been reported (Price and Drevets 2010). It is probable that the increase of glucose metabolism in this brain region is the compensatory mechanism acting at the initial stage of this disease and in unmedicated patients. In another research, an increased perfusion in the medial orbitofrontal cortex was observed during the induction of sadness in healthy controls (Liotti et al. 2002). Glucose hypermetabolism has also been used as a potential indicator of treatment response during the antidepressant therapy (Mayberg et al. 1997, 2000). Most frequently in antidepressant respondents, the increased dorsolateral and decreased ventrolateral prefrontal metabolism have been reported (Kennedy et al. 2007).

Interestingly, researchers have also found a correlation between the amygdala volume and glucose metabolism in the human brain in vivo (Johansen-Berg et al. 2008). Reduction in gray matter volume in the amygdala is probably compensated by an increased glucose metabolism at the beginning of a depressive episode. This prolonged compensation could provoke neuroplastic changes in the brain structure.

# 8 The Role of Glial Cells in Neurodegenerative Processes in Depression

Changes in the structure or function of neurons may be the result of disturbed action of glial cells (astrocytes, oligodendrocytes, and microglia). Astrocytes regulate many brain functions, including extracellular concentration of ions and neurotransmitters, glucose uptake, synthesis of neurotransmitters, synaptic function, neuronal metabolism, and migration. Astrocytes and microglia are also a source of many neurotrophic factors and cytokines.

Many studies have indicated the role of glial elements in the pathophysiology of various neurodegenerative disorders, including depression (Rajkowska and Miguel-Hidalgo 2007). It has been shown that glial ablation in the PFC evoked by an astrocyte-specific toxin, L-alpha-aminoadipic acid, can sufficiently induce depressive-like behaviors in rodents (Banasr and Duman 2008). Numerous studies reported a decrease of the glial density in various brain structures in depression. For instance, it has been revealed in layers 3 and 5 of the dorsolateral prefrontal cortex and in the deeper layers of the caudal orbitofrontal cortex (Rajkowska et al. 1999). Reduction of glial cell density was also observed in the subgenual part of the anterior cingulate cortex and in the supracallosal part of the anterior cingulate cortex in depressive subjects (Ongur et al. 1998; Cotter et al. 2001). By contrast, in hippocampal CA subfields and in the granule cell layer of the dentate gyrus, the glial cell density was increased in subjects with depression (Stockmeier et al. 2004).

Experimental data indicated that glucocorticoids, which appear to be elevated in depression, reduce the activation of astrocytes in the hippocampus (Sonino et al. 1993; Laping et al. 1994). According to this, concentration of glial fibrillary acidic protein (GFAP), a protein expressed in astrocytes, is also decreased in the hippocampus after chronic psychosocial stress (Czéh et al. 2006). Since a decreased number or function of astrocytes attenuates synaptic glutamate uptake and in the end enhances synaptic glutamate concentration, it may be one of the mechanisms responsible for neurodegenerative changes observed in depression. Glial cells are the main source of proinflammatory cytokines, which then regulate the kynurenine pathway, the main pathway for brain tryptophan metabolism (Schwarcz and Pellicciari 2002). Microglia cells, macrophages, and monocytes produce quinolinic acid, which is an agonist of the NMDA receptor, while astrocytes synthesize kynurenic acid, which is its antagonist (Guillemin et al. 2000, 2001, 2005). Increased quinolinic acid levels were found in blood and cerebrospinal fluid in major depression and in the cytokine-induced animal model of this disease (Myint et al. 2007; Dantzer et al. 2008; Raison et al. 2010), while higher kynurenic acid production was linked to schizophrenia (Erhardt et al. 2001; Nilsson et al. 2005; Linderholm et al. 2012). In this context, quinolinic acid may be the link between immune and neurotransmitter changes that are specific to depression and involved in neurodegenerative changes (Steiner et al. 2011). Moreover, glial cells are also the source and the target of various cytokines, e.g., interleukins, tumor necrosis factor, whose level appears to be elevated in depression (Müller and Ackenheil 1998; Howren et al. 2009; Dowlati et al. 2010).

An important function of astrocytes is glucose metabolism and providing energy to neurons. The majority of research with positron emission tomography and functional magnetic resonance, which is based on the activities of glucose uptake, glycolysis, and oxidative phosphorylation, indicated hypometabolism in the prefrontal cortex in depression patients. As such, these changes might be evoked by either a reduced number or weaker activation of astrocytes.

Properly functioning glial cells are necessary for the appropriate function of the neurons, including resistance to damage. It has been found that brain injury is associated with activation of astrocytes and manifested by the increase of the GFAP expression (Krohn et al. 1999). Activated astrocytes may promote neuronal survival by enhanced release of trophic factors but, conversely, they may also inhibit the process of regeneration by forming a glial scar.

Microglia, similar to astrocytes, regulate the extracellular concentration of ions and neurotransmitters and via the release of the trophic factor act on neuroprotective/ neurodegenerative processes. However, overstimulation of microglia seems to be harmful for brain cells. Microglia is activated mainly in response to immunity stimulation and brain damage, and this process is associated with the transition of microglia cells into the amoeboid form, which promotes phagocytosis and mobility (Streit et al. 1988; Kreutzberg 1996; Streit et al. 1999; Liu and Hong 2003). Activation of microglia, in response to proinflammatory triggers such as LPS (Pei et al. 2007; Qin et al. 2005a, b),  $\beta$ -amyloid (A $\beta$ ) (Bamberger and Landreth 2001), and HIV-Tat (Turchan-Cholewo et al. 2009), is associated with the liberation of cytotoxic agents, including TNF- $\alpha$  (Sawada et al. 1989; Lee et al. 1993), inflammatory prostaglandins (Wang et al. 2005), or reactive oxygen species (ROS) such as superoxide  $(O2^{\bullet-})$ (Colton and Gilbert 1987) and NO (Moss and Bates 2001; Liu et al. 2002), for the neutralization of pathogens (Oehmichen and Gencic 1975; Graeber et al. 1988). These compounds, besides the hydrogen peroxidase and hydroxyl radicals also produced by microglia, may injure neurons and glial cells (Chao et al. 1992; Boje and Arora 1992). Several studies revealed that antidepressant drugs (selective serotonin reuptake inhibitors, SSRI) modulate the ability of microglia to produce TNF- $\alpha$  and NO in mice (Hashioka et al. 2007, 2009; Hwang et al. 2008; Horikawa et al. 2010). Fluoxetine, sertraline, paroxetine, fluvoxamine, and citalopram have been reported to strongly inhibit the production of TNF- $\alpha$  and NO in LPS-stimulated microglial cells (Tynan et al. 2012). The above reports support the hypothesis of the role of activated microglia in the pathogenesis of depression. Furthermore, activation of microglia and an increased corticosterone level can appear together. In the restraint model of psychological stress in mice, the elevated corticosterone level was accompanied by an increased proliferation of microglia cells. The inhibition of glucocorticoid synthesis or antagonists of either the glucocorticoid receptor or NMDA receptor resulted in suppression of the stress-induced microglia cell proliferation (Nair and Bonneau 2006).

# 9 Effects of Antidepressant Drugs on Neurodegenerative Processes in Depression

Currently used antidepressants show a large variation in the mechanism of their action. Most registered drugs are based on the monoaminergic hypothesis of depression. According to this, the effect is acquired by increasing noradrenergic and/or serotonergic transmission through inhibition of monoamine's reuptake, the influence on adrenergic and serotonergic receptors, or inhibition of enzymatic degradation of monoamines. Tricyclic antidepressants, being the oldest group of drugs with a particularly broad spectrum of activity, have a number of adverse effects. The pharmacological action of antidepressants, as well as their toxicity, should be considered always in terms of two processes: changes after a single dose of the substance and the activity of the drug after prolonged use, due to changes in quantity/activity of the receptors, the intracellular molecular changes, and the process of plasticity. In this chapter, we focus on the effect and mechanism of action of antidepressant drugs on neuronal maintenance, the process of neurogenesis, and their influence on the morphological changes that occur during the state of depression. Although some antidepressant drugs exerted neuroprotective action on neurons and glial cells, some of these drugs, especially at higher doses, may exert a cytotoxic effect. For example, amitriptyline, desipramine, and fluoxetine induced in in vitro studies at µM concentrations cell death in hippocampal neurons and PC12 pheochromocytoma cells, and some antidepressants from the TCA and SSRI group induce apoptosis in neurons and glia cells (Španová et al. 1997; Post et al. 2000; Bartholomä et al. 2002).

### 9.1 Effects of Antidepressants on Proinflammatory Cytokines

The neuropharmacology of tricyclic antidepressants (TCA) cannot be described only in the example of noradrenaline (NA) and 5-HT reuptake inhibition because it has transpired that the profile of these drugs is much broader. The neuroprotective effect of tricyclic antidepressants has been described in the case of imipramine and clomipramine in in vitro studies. It has been shown that these drugs decrease the concentration of nitric oxide, TNF $\alpha$ , and IL-1 $\beta$ , as well as neurotoxic changes, observed in microglia cells and astrocytes stimulated with LPS (Hwang et al. 2008). This data is in line with the inflammatory and neurodegenerative theory of depression, whereby an increased level of proinflammatory cytokines and reactive oxygen species is purportedly involved in the pathogenesis of depression and the inhibition of microglia activation in the brain has a neuroprotective effect (e.g., Hwang et al. 2008; Maes et al. 2012). Similarly, it was reported that fluvoxamine, reboxetine, and imipramine inhibited the production of NO and IL-6 by microglial cells activated with IFN- $\gamma$  (Hashioka et al. 2007). In the in vivo experiment, it has been found that a 2-week administration of paroxetine inhibited, when stimulated by IFN $\alpha$ , IL-1 $\beta$  and IL-10 concentrations in the hypothalamus, while desipramine decreased the IL-1 $\beta$  and TNF $\alpha$  concentration in the rat cortex after LPS administration (Myint et al. 2007; O'Sullivan et al. 2009). When stimulated by LPS, the production of proinflammatory cytokines was also inhibited by an atypical antidepressant – tianeptine, but only in the hypothalamus and not in the hippocampus (Castanon et al. 2004). These studies indicated that antidepressant drugs showing a different action on monoamines inhibit the synthesis of proinflammatory cytokines.

# 9.2 Effects of Antidepressant Drugs on Neurogenesis

Recently, a lot of data has indicated that depression is associated not only with the process of neurodegeneration but also with reduced neurogenesis in the brain. Although the function of new neurons generated in the adult mammalian brain, especially in the dentate gyrus of the hippocampus, is poorly established, studies in animals suggest a relation between neurogenesis and some hippocampus-dependent learning task. Experimental data indicated that the administration of long-term antidepressants increases hippocampal neurogenesis and prevents the fall in this process induced by different factors, most often by various kinds of stress. For example, in chronic mild stress – an animal model of depression – a decrease in the total number of granule cells in the ventral hippocampus was observed, and long-term treatment with escitalopram blocked this reduction (Jayatissa et al. 2006). Also, in the olfactory bulbectomy model of depression, citalopram reversed inhibition of the new cells' creation in the dentate gyrus of the hippocampus (Jaako-Movits et al. 2006). Different classes of antidepressants (selective serotonin reuptake inhibitors, selective noradrenaline reuptake inhibitors, monoamine oxidase inhibitors, atypical antidepressants) and electroconvulsive seizures have been found to increase neurogenesis acting on the process of proliferation or survival of newborn neurons.

# 9.3 Effects of Antidepressant Drugs on the Brain Derived Neurotrophic Factor

The BDNF regulates the growth, survival, and function of brain cells and is involved in both the formation of memory traces and learning process. Some data has evidenced the involvement of BDNF in the pathogenesis of depression and in the action of antidepressant drugs. Chronic stress and glucocorticoids decreased BDNF expression in the hippocampus, whereas antidepressant drugs increase its concentration or prevent the stress effect. For example, it has been shown that in an animal model of chronic unpredictable stress, venlafaxine, mirtazapine, and fluoxetine raised the level of the BDNF in the hippocampus (Zhang et al. 2010). Also, desipramine, sertraline, and tianeptine, in some experimental models, increase the levels of BDNF. Chronic treatment with various antidepressant drugs also increases the expression or activity of TrkB (BDNF receptor), and BDNF/TrkB-signaling seems to be employed not only in the survival and differentiation of neurons but also in depression's etiology and the action of antidepressant drugs. It was revealed that in the TrkB knockout animals, antidepressant therapy has no effect, whereas application of BDNF into the hippocampus exerted an antidepressant-like effect (Shirayama et al. 2002; Saarelainen et al. 2003). In patients with depression, decreased levels of BDNF in the blood were normalized after long-term escitalopram treatment (Serra-Millàs et al. 2011). Also, postmortem studies showed decreased brain BDNF concentration in unmedicated depressed subjects (Pittenger and Duman 2008). A lot of data, but not all, indicated that BDNF is an important factor in the prevention of neuronal degeneration, the inhibition of neurogenesis, and may be essential for the action of antidepressants.

# 9.4 Effects of Antidepressant Drugs on the Atrophy of Hippocampal Neurons

Postmortem studies show that patients with depression have a significant atrophy of neuronal cells in the hippocampus (particularly area CA3 pyramidal neurons) and in the amygdala. In experimental animals, neurodegeneration is reversed after the application of antidepressant drugs belonging to different classes, such as fluoxetine, imipramine, tranylcypromine, rolipram, and reboxetine, plus also following electroconvulsive treatment or transcranial magnetic stimulation (Madsen et al. 2000; Mayberg et al. 2000; Czéh et al. 2002; Duman et al. 2004). In the inhibition of stressor glucocorticoid-induced hippocampal neurodegeneration, tianeptine especially exerts a protective function. This drug inhibits stress- or glucocorticoid-induced atrophy of apical dendrites of the hippocampal Ca3 pyramidal neurons (Magariños and McEwen 1995). Also, a stress-related decrease in the hippocampal volume, inhibition of cell proliferation, and a decreased level of N-acetylaspartate in the hippocampus and amygdala were prevented by tianeptine (Czéh et al. 2001; McEwen and Chattarji 2004). Tianeptine is an atypical antidepressant drug, which in contrast to other antidepressants increases serotonin uptake, and its neuroprotective action perhaps results from both the inhibition of NMDA receptor function and the increasing serotonin reuptake in the hippocampus, monoamine, whose release in the hippocampus is increased under many types of stress (Kole et al. 2002).

### 10 Zinc Deficiency-Induced Neurodegeneration in Depression

Recently, much attention has been paid to the role of zinc in the pathophysiology and treatment of depression (Szewczyk et al. 2010; Szewczyk et al. 2011b). The first strong evidence indicating hypozincemia in human depression was demonstrated by Maes and co-workers (Maes et al. 1994). On the other hand, zinc exhibits antidepressant activity in preclinical and clinical studies (Szewczyk et al. 2010, 2011a, b). Zinc deficiency is currently examined in experimentally induced depression in rodents (by a zinc deficient diet) and assessed in humans (e.g., Szewczyk et al. 2011a, b). Zinc deficiency induces depressive-like behavior and biochemical alterations similar to that observed in the animal models of depression (resembling human depression). The biochemical changes include reductions of brain BDNF, NGF concentrations, enhanced activity of the NMDA receptor, increased gluco-corticoid and glutamate actions, and reduced neurogenesis processes (e.g., Szewczyk et al. 2011a). Examinations of zinc deficiency as a model of depression are in progress.

### 11 Conclusion

Major depression is one of the most common psychiatric disorders whose etiology, despite long and intensive studies, remains unclear. Because about 30 % of patients do not respond to the therapy with currently available antidepressants, there is a great need to investigate new hypotheses explaining the cause of depression, plus create new drugs. Previous research on depression was focused on changes in the concentration of neurotransmitters and their receptors. Actually, increasing evidence indicates that an impaired neuronal plasticity, especially observed in the hippocampus and frontal cortex, plays a role in the pathogenesis of depression. In turn, the disruption of cellular mechanisms regulating neuronal plasticity is most often connected with prolonged, damaging effects of the glucocorticoids, glutamate, and proinflammatory cytokines. Thus, it seems that in the pathophysiology of depression, all of the main systems involved in homeostasis maintenance, namely, neurotransmitters, glucocorticoids, and cytokines, interact in a complex way. To study the role of one factor in the pathogenesis of depression is relatively easy, while the influence of several compounds and the determination of their interactions remain difficult. For example, even the interaction between glucocorticoids and proinflammatory cytokines is complex and depends on the duration of action and their actual concentration, plus involves the inhibitory action of glucocorticoids on cytokine synthesis, modulation of the cytokine receptor function, the stimulatory effect of some cytokines on cortisol synthesis, or their influence on glucocorticoid receptors. Moreover, it must be noted that the role of glucocorticoids, cytokines, and glutamate in the pathogenesis of depression has not yet been fully determined and that other factors may be also involved, while, furthermore, the pathogenesis of depression should not be connected only with cell neurodegeneration/death.

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Part IV

Neurotrophins

# Amyloid-Beta, BDNF, and the Mechanism of Neurodegeneration in Alzheimer's Disease

# Elyse Rosa and Margaret Fahnestock

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e-mail: rosae@mcmaster.ca; fahnest@mcmaster.ca

E. Rosa • M. Fahnestock (🖂)

Department of Psychiatry and Behavioural Neurosciences, McMaster University, Hamilton, ON, Canada

#### Abstract

The accumulation of amyloid- $\beta$  is widely considered the primary neurotoxic insult leading to Alzheimer's disease. Although the precise mechanism of this toxicity is not well understood, amyloid- $\beta$ -induced downregulation of brainderived neurotrophic factor may be one of the most important contributors to amyloid- $\beta$  toxicity. Brain-derived neurotrophic factor has diverse neurotrophic effects on the nervous system, including promoting synaptic plasticity and neurogenesis, and it is essential for cognition and memory. Early in the progression of Alzheimer's disease, prior to significant plaque and tangle deposition, declining brain-derived neurotrophic factor expression induced by amyloid- $\beta$  is associated with mounting impairments in cognition and memory. A variety of approaches have demonstrated that increasing brain-derived neurotrophic factor levels improves learning and memory, highlighting the possibility that therapeutically restoring brain-derived neurotrophic factor levels may prevent or reverse the memory impairments and cognitive decline seen in Alzheimer's disease.

#### Keywords

Alzheimer's disease • Amyloid- $\beta$  • BDNF • CREB • Learning and memory • LTP

### 1 Introduction

Alzheimer's disease (AD) is a progressive neurodegenerative disorder that is the most common cause of dementia in the elderly. AD presents as global cognitive decline with associated memory loss and altered personality. The neuropathological hallmarks of AD include extracellular amyloid- $\beta$  (A $\beta$ )-containing plaques, intracellular neurofibrillary tangles formed by hyperphosphorylated tau protein, synaptic loss, and resulting neurodegeneration (Hyman et al. 2012; Coleman & Yao 2003; Coleman and Flood 1987; Hyman 1997; Scheff and Price 2003; McKhann et al. 1984). AD primarily affects several regions of the brain that are known for their role in learning and memory, namely, the basal forebrain, hippocampus, cortex, and entorhinal cortex (Hyman et al. 1984; Coyle et al. 1983; Gomez-Isla et al. 1996).

Several theories exist to explain the molecular mechanisms that could lead to this devastating disease. One predominant theory suggests that an imbalance between A $\beta$  production and clearance produces the accumulation of A $\beta$  that is the driving force behind AD neuropathology (Hardy and Selkoe 2002; Iqbal and Grundke-Iqbal 2008). Amyloid- $\beta$  has many neurotoxic effects; however, the precise mechanism by which A $\beta$  results in the neurodegeneration seen in AD is unclear. The ability of A $\beta$  to downregulate the critical neurotrophic factor brainderived neurotrophic factor (BDNF) is the main focus of this chapter, as the promotion of synaptic integrity and function by BDNF is of utmost importance throughout life and for the prevention of learning and memory impairments.

### 2 Characteristics of Alzheimer's Disease

### 2.1 Pathological Features

Alzheimer's disease (AD) exhibits characteristic pathological features such as senile or amyloid plaques composed of precipitated A $\beta$  and neurofibrillary tangles composed of hyperphosphorylated tau; the presence of these pathologies is necessary for a definitive AD diagnosis (Hyman et al. 2012). The amyloid cascade hypothesis states that the accumulation of toxic A $\beta$  is the upstream driving force behind subsequent pathology and synaptic degeneration (Hardy and Selkoe 2002; Iqbal and Grundke-Iqbal 2008). Nevertheless, amyloid plaques do not correlate well with cognitive decline; neurofibrillary tangles correlate better (Guillozet et al. 2003). However, it is the loss of functional synapses in AD that correlates most strongly with loss of cognitive abilities (Terry et al. 1991).

A significant decline in synaptic connections in AD is followed by the pronounced loss of neurons in the basal forebrain, entorhinal cortex, hippocampus, and cortex. AD has been characterized as a disconnection of the hippocampus, as projections from the basal forebrain, entorhinal cortex, and cortex to the hippocampus are lost in AD (Hyman et al. 1984). Among these brain areas, the entorhinal cortex is one of the earliest areas vulnerable to synaptic degeneration, and because of its connection with the hippocampus and cortex, entorhinal cortex damage in AD significantly impacts learning and memory (Van Hoesen et al. 1991). In addition, studies have indicated that the memory deficits seen in AD are also directly related to the degree of basal forebrain cholinergic atrophy (Coyle et al. 1983). The basal forebrain is the major cholinergic output of the central nervous system, and these neurons as well as their projections to hippocampal and cortical regions are critical for learning, memory, and attention (Baxter and Chiba 1999). The loss of basal forebrain innervation is associated with aging and age-related memory loss (Terry and Katzman 2001; Ypsilanti et al. 2008). In fact, the most common drugs for AD on the market today are cholinergic enhancers.

In addition to loss of connectivity, there may be a loss of functional neurogenesis associated with AD. In many different AD transgenic mouse models, significantly less proliferation of new neurons is present in the hippocampal formation compared to control animals (Mu and Gage 2011). Without the generation of new neurons, the synaptic degeneration resulting from AD is not compensated and learning and memory are impaired. The case is not so clear in human hippocampus, however, where immunostaining with different markers of immature neurons suggests that there is either increased (Jin et al. 2004), no change (Boekhoorn et al. 2006), or decreased (Crews et al. 2010) neurogenesis in AD. More research is clearly needed to determine whether reduced hippocampal neurogenesis contributes significantly to memory loss in AD.

### 2.2 Clinical Presentation

#### 2.2.1 Impaired Cognition, Learning, and Memory

Perhaps the most clinically apparent symptom of the synaptic loss and neurodegeneration characteristic of AD is the decline of declarative memory, including both episodic and semantic memories (Aronoff et al. 2006). Initially, the loss of synaptic connections in the hippocampus and entorhinal cortex is evident as an early and significant decline in cognitive function and episodic memory (Devanand et al. 2007), which are hallmarks of AD. Episodic memory loss is not only an early event in AD, but in mild cognitive impairment (MCI), it is associated with an increased risk for conversion to AD (Aggarwal et al. 2005; Devanand et al. 2007). MCI is considered a preclinical stage of AD, since as many as 8 % of people with MCI progress to AD per year (Mitchell and Shiri-Feshki 2009). Transgenic mouse models of AD are also significantly impaired on tests of episodic-like memory compared to control animals (Davis et al. 2012). In addition, animal models of AD are impaired in long-term potentiation (LTP), a model for synaptic plasticity and a major cellular constituent underlying learning and memory processes. A $\beta$ , when injected into rat hippocampus or added to hippocampal slices, abolishes LTP without affecting baseline synaptic transmission (Cullen et al. 1997; Chen et al. 2000). Like episodic memory in humans, the LTP deficit in rodent models of AD occurs early; it occurs prior to plaque formation and is accompanied by deficits in spatial learning and memory (Liu et al. 2008; Jacobsen et al. 2006).

# 2.3 Molecular Pathway for Learning and Memory

#### 2.3.1 CREB

A molecular pathway essential for learning and memory that is disrupted in AD involves the transcription factor cAMP response element binding protein (CREB) (Barco et al. 2003). Aβ interferes with hippocampal LTP via signaling pathways including the Ca(2+)-dependent protein phosphatase calcineurin, Ca(2+)/calmodulindependent protein kinase II (CaMKII), and CREB (Yamin 2009). Active, phosphorylated CREB is reduced in the AD brain (Yamamoto-Sasaki et al. 1999) and is also decreased in neurons following treatment with A $\beta$  in vitro (Tong et al. 2001; Garzon and Fahnestock 2007). CREB phosphorylation recruits transcriptional coactivators which are required for the transcription of genes involved in learning and memory. Aβ-induced suppression of transcription induced by the CREB coactivator CRCT1 has been demonstrated in transgenic AD mice, and this suppression is mediated by blockade of L-type voltage-gated calcium channels, reduced calcium influx, and disruption of PP2B/calcineurin-dependent CRTC1 dephosphorylation (España et al. 2010). Restoration of CREB activity in AD transgenic mice by inducing expression of another coactivator, CREB binding protein (CBP), ameliorates learning and memory deficits (Caccamo et al. 2010). Among the genes induced by CREB and its coactivators that are required for learning and memory, the most important for AD may be brain-derived neurotrophic factor (BDNF).

# 3 BDNF Is a Powerful Neurotrophic Molecule with Pleiotropic Effects on the Nervous System

# 3.1 Neuronal Survival

BDNF supports the survival of neurons and their connections that are vulnerable in aging and in diseases of the aging brain such as AD, including the hippocampus, entorhinal cortex, neocortex, and basal forebrain (Hyman et al. 1984; Alderson et al. 1990: Knusel et al. 1991: Ghosh et al. 1994: Lindholm et al. 1996; Lowenstein and Arsenault 1996). Early studies demonstrated a more than twofold increase in survival of septal cholinergic neurons following BDNF treatment as measured by acetylcholinesterase (AChE) histochemical staining (Alderson et al. 1990). Further, blocking BDNF signaling either using antibodies or BDNF gene knockdown decreases survival of cortical (Ghosh et al. 1994). hippocampal (Lindholm et al. 1996), and dorsal root ganglion neurons (Korte et al. 1995). Although these results demonstrate that BDNF is a survivalpromoting factor for critical neuronal populations during development, conditional knockdown of BDNF in the adult suggests that BDNF is not required for postmitotic neuronal survival, but rather for dendritic maturation and growth of specific neuronal populations such as medium spiny neurons of the striatum (Rauskolb et al. 2010). Furthermore, the high-affinity BDNF receptor, tropomyosin-related kinase B (TrkB), unlike the related receptors TrkA or TrkC, is unable to trigger cell death in the absence of ligand (Nikoletopoulou et al. 2010), suggesting that decreases in BDNF may trigger altered spine morphology, reduced dendritic complexity, synaptic dysfunction, and degeneration but not outright neuronal loss.

# 3.2 Neurogenesis

BDNF promotes neurogenesis. Direct infusion of BDNF into rat hippocampus increases the number of developing neurons compared to saline injections in both the ipsilateral and contralateral hemispheres, suggesting a widespread effect (Scharfman et al. 2004). This hippocampal neurogenesis is directly related to subsequent cognitive functions, as new neurons are integrated into neural circuitry and play an important role in specific learning and memory tasks such as learning in the Morris water maze, a spatial memory task (Zhao et al. 2008).

Studies investigating the exact molecular mechanisms underlying BDNFmediated increases in neurogenesis are still underway; however, by signaling through the TrkB receptor, BDNF activates three signaling cascades, namely, the phosphatidyl-inosital-3-kinase (PI3K)/Akt, Ras/extracellular-regulated kinase (ERK), and phospholipase C gamma (PLC $\gamma$ ) pathways, all of which are implicated in BDNF's ability to activate CREB. Specifically, PLC $\gamma$  promotes the release of intracellular Ca(2+) stores (Berninger et al. 1993) which induce the activation of the CREB kinase, CaMKIV (Finkbeiner et al. 1997; Blanquet and Lamour 1997), and subsequently lead to the activating phosphorylation of CREB. Further, activation of the Ras/ERK signaling pathway is reportedly sufficient to promote Ca(+2)-/CRE-dependent transcription (Finkbeiner et al. 1997). Lastly, protein kinase A (PKA), which is activated by BDNF signaling predominantly via the PI3K/AKT pathway, can activate CREB by phosphorylating its activation site Ser133 (Gonzalez and Montminy 1989). In support of the latter pathway, the PKA activator rolipram increases the survival of new neurons in a CREB-dependent manner (Fujioka et al. 2004; Merz et al. 2012). The activation of CREB is associated with hippocampal neurogenesis, as phosphorylated CREB is consistently found in newborn hippocampal neurons (Nakagawa et al. 2002a, b; Jagasia et al. 2009). These findings support the possibility that BDNF promotes neurogenesis via mechanisms that rely on its ability to activate the neurogenesis-promoting transcription factor CREB.

### 3.3 Dendritic Modulation

BDNF regulates dendritic growth, branching and spine maturation through a balance between BDNF and its precursor form, proBDNF. Both mature and proBDNF signal through the TrkB receptor and the pan-neurotrophin receptor p75<sup>NTR</sup>; however, BDNF has a greater affinity for TrkB while proBDNF preferentially binds to p75<sup>NTR</sup> (Fayard et al. 2005; Teng et al. 2005). Signaling through TrkB promotes dendritic growth and spine maturation via activation of the PI3K/ Akt/mTOR signaling pathway, which has been long associated with axonal growth and dendritic complexity (Atwal et al. 2000; Kuruvilla et al. 2000; Markus et al. 2002; Kumar et al. 2005). Therefore, the preferential binding of BDNF to TrkB promotes synaptic development through increased dendritic growth and spine maturation. Conversely, proBDNF inhibits neurite outgrowth through its preferential binding to p75<sup>NTR</sup>, which negatively alters dendrite and spine morphology in addition to spine density (Yamashita et al. 1999; Zagrebelsky et al. 2005; Singh et al. 2008; Koshimizu et al. 2009). p75<sup>NTR</sup> reduces dendritic growth and branching through activation of a Rho-GTPase, RhoA (Sun et al. 2012), by releasing RhoA from its inactivating Rho-GDI (Yamashita and Tohyama 2003). Thus, the balance between proBDNF and BDNF plays a critical role in the regulation of spine maturation, dendritic growth/retraction, and synaptic density, which underlie learning and memory processes.

### 3.4 BDNF Plays a Critical Role in LTP

The cleavage of proBDNF to mature BDNF can take place extracellularly by plasmin or matrix metalloproteinases (MMPs) or intracellularly by furin or proprotein convertase 1 (PC1) (Mowla et al. 2001; Lu et al. 2005). The extracellular conversion by plasmin is specifically controlled by the extracellular serine protease tissue plasminogen activator (tPA) (Pang et al. 2004). tPA proteolytically cleaves inactive plasminogen to plasmin which can then cleave proBDNF into

mature BDNF (Lee et al. 2001). This conversion of plasminogen to plasmin by tPA takes place at hippocampal synapses, where subsequent conversion of proBDNF to mature BDNF occurs (Pang et al. 2004). Plasminogen-/- mice are impaired in late-phase long-term potentiation (L-LTP) (Pang et al. 2004). Addition of mature BDNF to these mice was able to rescue L-LTP, while addition of proBDNF was not, implicating proBDNF cleavage to mature BDNF in the mechanisms of LTP.

This ability of BDNF to rescue LTP is related to its preferential signaling through TrkB. Signaling through TrkB activates molecular cascades vital for synaptic integrity and for the production of LTP (Figurov et al. 1996). LTP is induced by the activation of protein kinases such as MAPK, PKA, PI3K, PKC, and CaMKII, which converge at the activation of ERK and modulate the activation of proteins involved in gene transcription and protein synthesis such as CREB (Lynch 2004). BDNF effectively activates protein kinases upstream of ERK through the PI3K/AKT, PKA, and Ras signaling pathways, which play a central role in the activation of CREB and subsequently contribute to neuronal survival and synaptic plasticity. CREB in turn plays a substantial role in upregulating the effector genes necessary for L-LTP (Pittenger and Kandel 1998) such as Arc, a member of the immediate-early gene (IEG) family that has been critically implicated in the maintenance of LTP (Ying et al. 2002; Guzowski et al. 2000), and BDNF itself. Additionally, BDNF is essential for maintaining the phenotype of VP16-CREB mice, which contain a constitutively active form of CREB and prime cells for L-LTP by activating CRE-mediated transcription (Barco et al. 2005). Conversely, proBDNF signaling through p75<sup>NTR</sup> favors long-term depression (LTD) (Lu 2003; Pang et al. 2004). Although the precise mechanism by which proBDNF signaling via p75<sup>NTR</sup> promotes LTD is still unknown, it may involve NMDA receptor subtype NR2B. NR2B activation is reportedly required for LTD (Massey et al. 2004), and proBDNF signaling through p75<sup>NTR</sup> directly potentiates the NR2B component of NMDA synaptic currents in hippocampal slices (Woo et al. 2005). Taken together, current research supports the model that regulation of BDNF/proBDNF processing modulates the balance of LTP/LTD, which underlies learning and memory.

# 3.5 BDNF Is Essential for Cognition and Memory

Multiple studies have shown that decreases in BDNF such as in heterozygous BDNF knockout mice or BDNF knockdown result in learning and memory deficits (Linnarsson et al. 1997; Gorski et al. 2003; Heldt et al. 2007). A Val66Met polymorphism in the pro-domain of proBDNF which interferes with BDNF secretion results in episodic memory impairments and reduced hippocampal volume, key endophenotypes of AD (Egan et al. 2003; Hariri et al. 2003). This finding shows that an alteration in BDNF levels is sufficient to affect hippocampal activity and that BDNF is critically involved in hippocampal learning and memory processes.

# 4 BDNF Is Significantly Downregulated in Alzheimer's Disease

### 4.1 BDNF Loss Is an Early Event in the Progression of AD

Amyloid plaques correlate poorly with cognition, and evidence suggests that cognitive decline precedes A $\beta$  pathology (Van Dam et al. 2003; Westerman et al. 2002; Hanna et al. 2009). This is supported by transgenic mouse models of AD, where deficits in novel object recognition and spontaneous alternation performance in a "Y" maze occur prior to the deposition of A $\beta$  plaques (Holcomb et al. 1998; Francis et al. 2012). Loss of BDNF also occurs prior to plaque deposition in these transgenic mice and coincides with memory deficits (Francis et al. 2012). Further, BDNF is lost early in the progression of AD, as demonstrated by a significant decline in BDNF in individuals with MCI. More than a 30 % reduction in cortical BDNF expression was found in MCI subjects compared to age- and gender-matched, noncognitively impaired controls (Peng et al. 2005), with expression levels nearly as low as in AD subjects. Lastly, like humans, canines develop cognitive impairment and amyloid deposition as they age and are an excellent model of MCI (Cotman & Berchtold 2002; Cotman & Head 2008). BDNF mRNA levels were significantly decreased and correlated with cognitive status in aged, cognitively impaired canines as well, supporting human findings that BDNF loss coincides with memory impairments (Fahnestock et al. 2012). Thus, in both animal models and humans, BDNF is lost early, coinciding with memory loss and prior to the appearance of significant plaque deposition typical of AD.

# 4.2 BDNF Is Indicative of Cognitive Status

The expression of BDNF is strongly correlated with cognitive status. BDNF mRNA and proBDNF and mature BDNF protein are decreased in the entorhinal, frontal, temporal, and parietal cortices as well as the hippocampus of AD subjects compared to controls (Ferrer et al. 1999; Phillips et al. 1991; Murray et al. 1994; Narisawa-Saito et al. 1996; Connor et al. 1997; Holsinger et al. 2000; Hock et al. 2000; Fahnestock et al. 2002; Michalski and Fahnestock 2003; Peng et al. 2005). Lower levels of both mature BDNF and proBDNF protein correlate with decreased cognitive test scores as measured by the Mini Mental State Examination (MMSE) and Global Cognitive Score (GCS) (Peng et al. 2005).

# 4.3 Increasing BDNF Levels Can Counteract Memory Impairments

The strong correlation between lowered BDNF and diminished cognition, learning, and memory makes the possibility of rescuing these behavioral symptoms of AD with BDNF an enticing undertaking. Knocking down BDNF in mice results in loss

of synapses, learning and memory deficits, and diminished LTP, whereas BDNF administration rescues or increases LTP and synapses and restores learning and memory (Blurton-Jones et al. 2009; Nagahara et al. 2009; Patterson et al. 1996; Korte et al. 1995). Early studies substantiated findings that BDNF is necessary for LTP: mice with a deletion in the BDNF gene exhibited significantly weakened hippocampal LTP expression (Korte et al. 1995), whereas treating hippocampal BDNF knockout tissue with recombinant BDNF completely restored LTP (Patterson et al. 1996). Similarly, knockdown of the BDNF receptor TrkB can significantly reduce hippocampal LTP (Minichiello et al. 1999). These early studies led to efforts in recent years to deliver exogenous BDNF as a method of restoring cognitive abilities in animal models of AD. However, expression of truncated TrkB receptors in the ventricular ependyma effectively prevents diffusion of BDNF to target tissues when it is administered by intracerebroventricular infusion (Anderson et al. 1995). Furthermore, BDNF does not readily cross the blood-brain barrier. Fusions of BDNF with factors that allow it to be transferred across the blood-brain barrier such as an antibody to the transferrin receptor (Zhang and Pardridge 2001a, b; 2006) or to the insulin receptor (Boado et al. 2007) have facilitated entry of intravenously administered BDNF into brain tissue.

The brain is exquisitely sensitive to BDNF levels, which presents another significant barrier to the successful use of this trophic factor in vivo. Microinjections of microgram levels of BDNF into the hippocampus increase neuronal excitability and can even cause seizures, whereas chronic infusions of similar amounts decrease the TrkB receptor and TrkB activation, resulting in loss of responsiveness to BDNF (Scharfman 1997; Xu et al. 2004). Furthermore, delivery of BDNF can result in impairment of spatial learning rather than improvement if BDNF expression is too high (Pietropaolo et al. 2007). Therefore, more physiological delivery of low doses of BDNF via viral vectors has been extensively investigated.

In the entorhinal cortex-lesioned aged rat, BDNF gene delivery enhanced LTP and partially restored cognitive function (Ando et al. 2002). BDNF gene delivery via lentiviral injection to the entorhinal cortex in a transgenic mouse model of AD not only increased synaptic density (synaptophysin-IR) and synaptic function (p-Erk signaling) but, importantly, improved spatial memory performance in the Morris water maze task (Nagahara et al. 2009). BDNF treatment did not affect neuronal number in this model, although it did rescue neurons from cell loss in a perforant path transection model. BDNF administration also exerted similar effects and rescued learning and memory in the aged, cognitively impaired nonhuman primate. Importantly, this lentiviral expression system has been used to maintain delivery of NGF for up to 1 year in primate basal forebrain, eliminating transient expression as a disadvantage of the system. These experiments were carried out after the onset of disease symptoms and pathology in transgenic mice and after cognitive impairment appeared in aged primates, suggesting that BDNF may be an effective therapy for reversal of cognitive deficits as well as prevention.

However, this delivery method suffers from poor viral diffusion within the parenchyma. This complication is somewhat ameliorated by the implantation of neural stem cells (NSC) which can migrate within brain parenchyma. NSC delivery

into hippocampus of the triple-transgenic (Tg3x-AD) mouse model of AD increased synaptic density and improved hippocampal-dependent learning and memory (Blurton-Jones et al. 2009). BDNF knockdown in these NSCs eliminated the beneficial effects of the cells, demonstrating that BDNF secreted by the NSCs was the active factor (Blurton-Jones et al. 2009).

Alternative approaches to gene delivery include methods of increasing endogenous BDNF levels or signaling. For instance, viral delivery of CREB activators (Caccamo et al. 2010; Espana et al. 2010), successfully reverses synaptic atrophy and learning and memory impairments in transgenic mice. Rolipram, a phosphodiesterase inhibitor and CREB activator, increases BDNF expression (DeMarch et al. 2008) and therefore may have therapeutic effects in AD. Additionally, small molecule BDNF mimetics or TrkB agonists which promote signaling through the TrkB receptor can restore synaptic plasticity and cognitive function (Massa et al. 2010). Moreover, natural products and their derivatives such as certain flavonoids are under active investigation as potential neuroprotective agents, LTP enhancers and activators of CREB, TrkB, and its downstream signaling pathways PI3K/Akt and MAPK/Erk (Maher et al. 2006; Jang et al. 2010). Finally, lifestyle adjustments such as exercise, environmental enrichment, and dietary restriction increase BDNF expression in both animal models and human studies (Mattson et al. 2004b). An enriched environment has been shown to increase BDNF mRNA in rats (Falkenberg et al. 1992), while an exercise routine can regulate transcript-specific BDNF mRNA and protein (Berchtold et al. 2002; Cotman and Berchtold 2002; Zajac et al. 2010). Recently, the combination of an antioxidant-rich diet and environmental enrichment was shown to significantly increase BDNF expression and was correlated with an improvement in memory scores in old, cognitively impaired canines (Fahnestock et al. 2012). Although a variety of molecular, pharmacological, and lifestyle interventions are under investigation and may pave the way for future clinical trials in subjects with AD, increasing brain BDNF levels safely and effectively is not yet a reality. A greater understanding of BDNF regulation and expression may provide additional targets or modalities.

# 4.4 BDNF Transcriptional and Posttranslational Regulation

The human BDNF gene is 70 kb long and consists of 11 exons and 9 functional promoters, which by alternative splicing result in at least 17 different BDNF transcripts (Pruunsild et al. 2007). This variation in transcript expression allows for tissue-specific regulation of BDNF in response to a variety of developmental and environmental cues (Timmusk et al. 1993; Pruunsild et al. 2007). BDNF transcripts II, III, IV, V, and VII are exclusively localized to the brain, while other BDNF transcripts are expressed in neuronal as well as nonneuronal tissues (Pruunsild et al. 2007). Within the brain, BDNF transcript IV accounts for approximately half of the total BDNF mRNA found in the cortex (Garzon and Fahnestock 2007). BDNF transcript IV is regulated at least in part by CREB (Shieh et al. 1998; Pruunsild et al. 2007; Timmusk et al. 1995).

BDNF protein is synthesized as 36 kDa precursor proBDNF, which can be cleaved by enzymes including plasmin, furin, and matrix metalloproteases to produce 14 kDa mature BDNF (Mowla et al. 2001; Lee et al. 2001; Lu et al. 2005). ProBDNF is expressed widely in the human brain, where it is only partially cleaved to its mature form. Both BDNF and proBDNF are found in relatively high amounts in the hippocampus, cortex, and basal forebrain (Michalski and Fahnestock 2003), areas that are particularly vulnerable in AD.

### 4.5 BDNF mRNA and Protein Decrease in AD

BDNF levels are dramatically altered in individuals with AD and MCI. BDNF mRNA expression in the hippocampus is decreased nearly twofold in individuals with AD (Phillips et al. 1991), and BDNF mRNA is reduced approximately threefold in the AD parietal cortex (Garzon et al. 2002; Holsinger et al. 2000). This decrease in BDNF mRNA is due to a decrease in four BDNF transcripts: I, II, IV, and VI (Garzon et al. 2002). Of particular interest is transcript IV, since it is the most prevalent transcript in human cortex (Garzon and Fahnestock 2007) and is specifically decreased in human and mouse models of AD (Garzon et al. 2002; Peng et al. 2009).

The decreased BDNF mRNA in AD corresponds to a similar decrease in BDNF protein. Mature BDNF protein is significantly reduced in AD hippocampus (Narisawa-Saito et al. 1996; Connor et al. 1997; Hock et al. 2000), frontal cortex (Ferrer et al. 1999), temporal cortex (Connor et al. 1997), and parietal cortex (Hock et al. 1998; Peng et al. 2005) compared to controls, and proBDNF protein is reduced in AD parietal cortex (Michalski and Fahnestock 2003; Peng et al. 2005). The molecular mechanisms in AD that could lead to this substantial decrease in BDNF expression are beginning to be understood.

### 5 Amyloid-β Toxicity in Alzheimer's Disease

Although A $\beta$  has been recognized as having non-disease associations, such as acting as an antimicrobial agent (Soscia et al. 2010), it is most well studied as the precipitating insult in AD. Early studies suggested that it was the formation of A $\beta$  plaques that was the primary neurotoxic insult in Alzheimer's disease. However, plaques do not correlate well with decreased cognition, and more recent evidence suggests that soluble forms of A $\beta$  may be more toxic (Ferreira et al. 2007; Walsh and Selkoe 2007; Lacor et al. 2007; Hardy and Selkoe 2002; Garzon and Fahnestock 2007).

### 5.1 Formation of Toxic, Soluble A $\beta$

Amyloid- $\beta$  is a peptide formed by the proteolytic processing of the amyloid precursor protein (APP) (Hardy and Selkoe 2002; Haass et al. 1992; Citron et al.

1992; Shoji et al. 1992). APP is a single-pass, transmembrane protein with a small intracellular C-terminus and a large extracellular segment which contains part of the A $\beta$  sequence (Mattson 2004). APP is initially proteolytically cleaved by  $\alpha$ - or  $\beta$ -secretase (also known as beta-site APP cleaving enzyme 1; BACE-1), which releases a soluble fragment of APP (DeStrooper and Annaert 2000). Following primary cleavage by  $\beta$ - or  $\alpha$ -secretase, a secondary cleavage of the transmembrane domain, or C-terminal fragment, of APP by y-secretase results in the formation of either the A $\beta$  peptide or the p3 fragment of APP, respectively (DeStrooper and Annaert 2000; Haass et al. 1992). Gamma-secretase is a complex of integral membrane proteins that includes presenilin-1 (PS-1) or presenilin-2 (PS-2), which when mutated can dramatically alter the production of A $\beta$ . Gamma-secretase cleavage following cleavage by  $\beta$ -secretase can result in A $\beta$  peptides of various lengths (Haass et al. 1992; DeStrooper and Annaert 2000).  $A\beta_{1-40}$  is most commonly produced following the successive cleavage of  $\beta$ - and  $\gamma$ -secretases; however, it is  $A\beta_{1-42}$  that is of particular interest in the progression of AD because of its greater propensity to aggregate (Tabaton et al. 2010; Glabe 2001). The aggregation of A $\beta$  involves several conformational states from dimers and trimers to high-molecular-weight oligomers and protofibrils and finally insoluble A $\beta$  fibrils (Glabe 2004) and requires a structural change from an  $\alpha$ -helical conformation to a more organized  $\beta$ -sheet configuration characteristic of A $\beta$  aggregation (Xu et al. 2005).

Formation of soluble A $\beta$  is increased by mutations in APP, PS-1, and PS-2, hallmarks of familial forms of AD (Citron et al. 1992; Haass et al. 1994; Hutton and Hardy 1997; Scheuner et al. 1996). Such mutations result either in A $\beta$  overproduction or in an increase in the aggregation-prone A $\beta_{1-42}$  (Burdick et al. 1992; Jarrett et al. 1993; Hardy and Selkoe 2002; Selkoe 1994). Increased levels of A $\beta$  increase aggregation and toxicity prior to plaque deposition. Levels of soluble A $\beta$  oligomers in the frontal cortex of individuals with AD are up to 70-fold higher than in control brains (Gong et al. 2003) and correlate with the degree of dementia and of synaptic loss (Lue et al. 1999).

# 5.2 Mechanisms of Aβ Toxicity

A $\beta$  has many proposed toxic effects that could result in the neuronal dysfunction and neurodegeneration characteristic of AD. A $\beta$  enhances tau hyperphosphorylation and the subsequent formation of neurofibrillary tangles in animal models (Gotz et al. 2001; Lewis et al. 2001; Masliah et al. 2001; Hardy and Selkoe 2002) and in human neuroblastoma cells in vitro (Pennanen and Gotz 2005). This suggests that A $\beta$  may exert its neurotoxic effects via tau hyperphosphorylation and subsequent neurofibrillary tangle formation (Iqbal and Grundke-Iqbal 2008). Another well-studied toxic effect of A $\beta$  accumulation is altered calcium homeostasis: soluble A $\beta$  oligomers have been shown to alter calcium regulation by modulating ion channels, including voltage-gated calcium and potassium channels and nicotinic and NMDA receptors, and by forming its own calcium-conducting

pores, thereby increasing levels of cytosolic calcium (Green et al. 2007; Mattson et al. 1993; Ferreiro et al. 2006; Demuro et al. 2005; Resende et al. 2007). Oligomeric A $\beta$  has also been shown to promote increased reactive oxygen species (ROS) production in primary cortical and hippocampal neurons (Sponne et al. 2003; DeFelice et al. 2007), which has been associated with inflammation and neurodegeneration. Amyloid- $\beta$  may also indirectly induce neurotoxicity through the activation of proinflammatory responses (Akiyama et al. 2000; Giovannini et al. 2002; Lukiw and Bazan 2000), which may have a role in inhibiting hippocampal memory formation (Tancredi et al. 1992, 2000; Murray and Lynch 1998; Heneka and O'Banion 2007). However, A $\beta$  has its most direct effect on synaptic degeneration through its action on specific signaling cascades.

# **6 BDNF May Mediate A**β Toxicity in Alzheimer's Disease

A $\beta$  toxicity results in neurodegeneration by altering signaling pathways upstream and downstream of CREB. CREB phosphorylation and signaling are reduced by exposure to soluble, oligomeric A $\beta$  in vitro (Tong et al. 2001; Garzon and Fahnestock 2007). A $\beta$  treatment inhibits the Ras/ERK and PI3K/AKT signaling pathways (Tong et al. 2004), which are downstream of BDNF/TrkB but are also essential for the phosphorylation of CREB and subsequent expression of BDNF. It has recently been shown that inhibiting glycogen synthase-3 $\beta$  (GSK3 $\beta$ ), which can be activated by A $\beta$ , is an effective way to decrease inhibitory phosphorylation of CREB and increase BDNF protein (DaRocha-Souto et al. 2012). Additionally, A $\beta$ has been shown to inactivate PKA in vitro, and increasing PKA activity reverses decreased CREB activation induced by A $\beta$  treatment (Vitolo et al. 2002). Therefore, A $\beta$  disrupts several kinase cascades, leading to a reduction in activity of CREB and in BDNF expression.

Soluble, oligomeric A $\beta$ , but not the fibrillar A $\beta$  found in plaques, decreases CREB activation and downregulates BDNF in vitro, largely via transcript IV (Garzon and Fahnestock 2007). In transgenic mouse models of AD, which also exhibit downregulation of transcript IV, BDNF expression is inversely proportional to the amount of soluble, high-molecular-weight A $\beta$  oligomers (Peng et al. 2009). Thus, inactivation of CREB by soluble, aggregated A $\beta$  may downregulate BDNF transcript IV, resulting in the diminished synaptic connections and memory loss characteristic of AD.

A major consequence of  $A\beta$ -induced CREB reduction that is mediated by BDNF is diminished LTP. Soluble  $A\beta$  oligomers have been shown to effectively inhibit hippocampal LTP (Lambert et al. 1998; Walsh et al. 2002).  $A\beta$  inhibits LTP by specifically interfering with CaMKII and CREB signaling pathways (Yamin 2009), and  $A\beta$ -induced reductions in LTP are rescued by BDNF (Zeng et al. 2010). These effects on LTP provide a mechanism for  $A\beta$ 's interference with learning and memory. Therefore, although  $A\beta$  has many proposed mechanisms of toxicity, its modulation of kinase pathways affecting transcription of genes involved in learning and memory, particularly BDNF and CREB, is perhaps the most significant for AD.

### 7 Conclusion

Alzheimer's disease, which increasingly affects our aging population, is primarily a disease of synaptic dysfunction. Soluble, aggregated amyloid- $\beta$  is thought to be the primary neurotoxic insult leading to synaptic loss. However, the exact mechanisms that lead from A $\beta$  aggregation to the pathological and physical symptoms of AD are not clear. A $\beta$  accumulation, among other toxic effects, alters kinase cascades, leading to reduced CREB activation and to BDNF downregulation, primarily through transcript IV. This downregulation of BDNF is an early damaging event in the progression of AD and occurs in MCI and prior to plaque deposition in transgenic mice. BDNF promotes the integrity and function of synaptic connections and the function of neuronal circuits required for cognition, learning, and memory, and it is one of the most powerful neural repair molecules known. BDNF administration in animal models of AD can reverse morphological and behavioral symptoms even when administered late in the disease process. This suggests that the therapeutic use of BDNF might arrest or even correct the cognitive decline seen so early in the progression of AD. Methods to deliver or enhance BDNF and its signaling cascade are under investigation. A deeper understanding of the mechanisms that regulate BDNF in AD may provide additional therapeutic options.

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# BDNF V66M Polymorphism and Brain Functions

# Zhe-Yu Chen

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#### Abstract

Brain derived neurotrophic factor (BDNF) plays important roles in neuronal survival, axonal and dendritic growth, synaptic plasticity, learning and memory and emotional behavior. Recently, a single nucleotide polymorphism (SNP) in the BDNF gene leading to a valine to methionine substitution at position 66 in BDNF prodomain (Val66Met) has been found to decrease activity-dependent BDNF secretion and to be associated with increased susceptibility to neuropsychiatric disorders. BDNF Val66Met polymorphism only exists in human and represents the first genetic alteration in a neurotrophin that has been linked to a human disease. In this chapter, we will focus on the biological consequences of BDNFVal66Met polymorphism in the context of central nervous system structure and function with combined molecular and cellular, human behavioral and imaging, and transgenic mouse studies.

#### Keywords

 $BDNF_{Met}$   $\bullet$  Behavior  $\bullet$  Brain structure  $\bullet$  Neuropsychiatric disorders  $\bullet$  Secretion

#### Z.-Y. Chen

Department of Neurobiology, School of Medicine, Shandong University, Jinan, Shandong, China e-mail: <a href="mailto:zheyuchen@sdu.edu.cn">zheyuchen@sdu.edu.cn</a>

### 1 Introduction

Brain-derived neurotrophic factor (BDNF), a molecule known to regulate neuronal survival and differentiation, plays an important role in activity-dependent plasticity processes, such as long-term potentiation (LTP), learning, and memory (Huang and Reichardt 2001; Chao 2003). BDNF is widely expressed in the development and adult mammalian brain and has been implicated in the actions of both antidepressants and mood stabilizers, as well as the pathophysiology of affective disorders (Schuman 1999; Castren and Rantamaki 2010; Krishnan and Nestler 2010). Therefore, BDNF plays a significant role in regulating specific synaptic functioning in learning, memory, and neuropsychiatric disorders. Recently, a single nucleotide polymorphism (SNP) in the BDNF gene leading to a valine-to-methionine substitution at position 66 in BDNF prodomain (Val66Met) has been identified and shown to influence human hippocampal volume and memory (Egan et al. 2003). This BDNF polymorphism is found to be associated with altered susceptibility to neuropsychiatric disorders including bipolar disorder, Alzheimer's disease, and depression (Neves-Pereira et al. 2002; Sklar et al. 2002; Ventriglia et al. 2002; Nederhof et al. 2010). BDNF Val66Met polymorphism only exists in human and represents the first genetic alteration in a neurotrophin that has been linked to a human disease. This chapter will be focused on the biological consequences of BDNF Val66Met polymorphism in the context of central nervous system structure and function.

# 2 BDNF<sub>Met</sub> SNP and Mental Disorders

The human BDNF gene is located on the short arm of chromosome 11 (11p14.1). One frequent, nonconservative polymorphism in the human BDNF gene (dbSNP number rs6265) has been identified, a single nucleotide polymorphism (SNP) at nucleotide 196 (G/A) producing an amino acid substitution (valine to methionine) at codon 66 (Val66Met). The frequencies for the alleles of the Val66Met polymorphism were A = 20–30 % and G = 70–80 % in Caucasians (Neves-Pereira et al. 2002; Egan et al. 2003). However, the frequency of BDNF<sub>Met</sub> SNP varies depending upon ethnicity. The frequency of healthy individuals who carried G/G (Val/Val) was significantly decreased in Japanese (33.8 %) than that in Italians (48.7 %) or in Americans (68.4 %) (Shimizu et al. 2004), which suggests that BDNF might act to produce ethnic traits.

Accumulating evidence suggests that BDNF plays a role in the pathophysiology of psychiatric diseases. A recent series of studies have linked BDNF Val66Met polymorphism with memory impairments as well as altered susceptibility to neuropsychiatric disorders, such as bipolar disorders, Alzheimer's disease, Parkinson's disease, depression, and eating disorders (Neves-Pereira et al. 2002; Ventriglia et al. 2002; Masaki et al. 2003; Ribases et al. 2003; Schumacher et al. 2005). This BDNF polymorphism represents the first alternation in a neurotrophin gene that has been linked to clinical pathology. Anxiety is a common symptom among most psychiatric disorders. Several recent studies have looked at the relationship between the BDNF<sub>Met</sub> variant and the anxiety traits. The results have been conflicting, with the Val allele associated with vulnerability in one study and the Met allele designated as the "risk" allele in another study (Sen et al. 2003; Jiang et al. 2005; Lang et al. 2005). Inconsistency across genetic studies may be attributable to sampling and measurement issues, genetic heterogeneity due to differential sampling of populations, or low frequency of homozygous Met carriers, which may lessen the effect size of any particular association. It may also relate to a failure to take into account relevant geneby-gene and gene-by-environment interactions. A recent investigation found the association between incident stroke and depression with the strongest association for Met/Met genotype participants (Kim et al. 2008), which provides evidence for a gene-environment interaction with respect to the impact of stroke on depression. Other studies revealed that BDNF interacted with childhood adversity or stressful life events that predicted greater likelihood of depression among Met carriers and did not influence Val/Val carriers, which further confirmed that the association of BDNF<sub>Met</sub> SNP with depression was modified by the environment factor such as stress (Bukh et al. 2009; Carver et al. 2011). A significant three-way interaction between BDNF genotype, 5-HTTLPR, and maltreatment history in predicting depression was also reported (Kaufman et al. 2006). Children with the Met allele of the BDNF gene and two short alleles of 5-HTTLPR had the highest depression scores, but the vulnerability associated with these two genotypes was only evident in the maltreated children. To further elucidate the role of BDNF<sub>Met</sub> variant in psychiatric disease, the BDNF<sub>Met</sub> knock-in mouse would be an ideal tool since the transgenic mouse would provide a more homogeneous gene background and the gene-by-gene and gene-by-environment effects will be more easily controlled in mouse behavior study.

## 3 Molecular Mechanism Underlying BDNF<sub>Met</sub> Variant

Brain-derived neurotrophic factor (BDNF) has been shown to play significant roles in neuronal survival, axonal and dendritic growth and guidance, synaptic structure and connections, neurotransmitter release, LTP, and synaptic plasticity (Chao 2003). Thus, the influence of BDNF spans from developmental neurobiology to neurodegenerative and psychiatric disorders. The actions of BDNF are dictated by two classes of cell surface receptors, the TrkB receptor tyrosine kinase and the p75 neurotrophin receptor (p75<sup>NTR</sup>), a member of the TNF receptor superfamily (Chao et al. 2006). BDNF binding to TrkB receptor triggers receptor dimerization, transphosphorylation of intracellular tyrosine residues, and subsequent activation of the PI-3 kinase, Ras/MAPK, and PLC- $\gamma$  pathways, thus influencing transcriptional events that have a multitude of effects on cell cycle, neurite outgrowth, and synaptic plasticity (Huang and Reichardt 2003; Teng and Hempstead 2004; Chao et al. 2006). Signal transduction through p75 independently gives rise to increase in JNK (c-Jun N-terminal kinase), NF- $\kappa$ B (nuclear factor  $\kappa$ B), and ceramide, thus triggering apoptosis (Roux and Barker 2002). Consistent with the critical role of BDNF in synaptic plasticity, BDNF is synthesized and released in a manner that is dependent on neuronal activity (Lessmann et al. 2003; Lu 2003; Bramham and Messaoudi 2005). In the mammalian brain, BDNF is synthesized as a precursor called proBDNF, which is proteolytically cleaved to generate mature BDNF. ProBDNF may preferentially bind p75<sup>NTR</sup>, whereas mature BDNF preferentially binds TrkB receptor (Teng et al. 2005; Nagappan et al. 2009; Yang et al. 2009).

The molecular mechanisms underlying altered  $BDNF_{Met}$  function have begun to be studied primarily in vitro cell culture systems. The Met substitution in BDNF prodomain was shown in neurosecretory cells and primary cultured neurons to lead to four trafficking defects: (1) decreased variant  $BDNF_{Met}$  distribution into neuronal dendrites, (2) decreased variant  $BDNF_{Met}$  targeting to secretory granules, (3) subsequent impairment in BDNF-regulated secretion, and (4) impaired dendritic targeting of BDNF mRNA (Egan et al. 2003; Chen et al. 2004; Chen et al. 2005; Chiaruttini et al. 2009). The molecular mechanisms underlying the trafficking defects associated with  $BDNF_{Met}$  variant are described in detail below.

When expressed together in the same cell, BDNF<sub>Met</sub> alters the trafficking of wild-type BDNF (BDNF<sub>Val</sub>) through the formation of heterodimers that are less efficiently sorted into the regulated secretory pathway (Chen et al. 2004). These findings are consistent with previous studies indicating that the prodomain of neurotrophins plays important role in regulating their intracellular trafficking to secretory pathways (Suter et al. 1991; Mowla et al. 1999). These in vitro studies with BDNF<sub>Met</sub> point to the presence of a specific trafficking signal in the BDNF prodomain region encompassing the Met substitution that is required for efficient BDNF sorting. Following this hypothesis, a trafficking protein, sortilin, was identified to be necessary for the efficient sorting of BDNF to the regulated secretory pathway. Sortilin interacts specifically with BDNF in a region encompassing the Met substitution (Chen et al. 2005). Replacement of Met at this position led to decreased interaction of BDNF with sortilin and suggests that decreased protein-protein interaction between BDNF and the trafficking machinery is one plausible molecular model for the secretion defect observed with the variant BDNF. Targeting of BDNF mRNA to dendrites also plays a key role in mediating synaptic plasticity (An et al. 2008; Tongiorgi and Baj 2008). Constitutive dendritic targeting of BDNF mRNA is mediated by translin, an RNA-binding protein implicated in RNA trafficking (Chiaruttini et al. 2009). BDNF<sub>Met</sub> mutation blocks dendritic targeting of BDNF mRNA by disrupting its interaction with translin (Chiaruttini et al. 2009). A BDNF<sub>Met</sub> knock-in transgenic mouse was recently generated and provided a unique tool to study the in vivo consequences of BDNF<sub>Met</sub> SNP (Chen et al. 2006). The transcription of BDNF<sub>Met</sub> is regulated by endogenous BDNF promoters in the BDNF<sub>Met</sub> knock-in mouse, which fully mimics the human BDNF<sub>Met</sub> polymorphism. When hippocampal neurons were cultured from BDNF<sub>Met</sub> mice, a significant decrease in the BDNF-regulated secretion was detected, which suggested an insufficiency in available BDNF in BDNF<sub>Met</sub> mice (Chen et al. 2006). In BDNF<sub>Met/Met</sub> mice, pilocarpine-induced dendritic trafficking

of BDNF mRNA was also blocked, which further confirmed the in vitro study (Chiaruttini et al. 2009). Thus, the in vitro findings of the trafficking defects associated with  $BDNF_{Met}$  variant were verified in  $BDNF_{Met}$  knock-in mouse.

It also remains possible that there are additional defects in  $BDNF_{Met}$  processing, although in vitro studies in neurons suggest no defect in  $BDNF_{Met}$  processing (Egan et al. 2003; Chen et al. 2004). It has been reported that tissue plasminogen activator, by activating the extracellular protease plasmin, converts proBDNF to the mature BDNF and that such conversion is critical for late-phase long-term potentiation expression in the mouse hippocampus (Pang et al. 2004). Given that proBDNF preferentially activates p75<sup>NTR</sup> over TrkB receptor, it is likely that proteolytic conversation of proBDNF may be implicated in the BDNF function. Whether BDNF<sub>Met</sub> affects BDNF processing should be further examined in vivo in the BDNF<sub>Met</sub> knock-in mice.

# 4 Brain Structure Alterations Associated with BDNF<sub>Met</sub> Polymorphism

The fundamental question is what the in vivo consequences of  $BDNF_{Met}$  SNP are. One of the most reliable effects observed in carriers of the Met allele is a reduction in hippocampal volume. In studies of human brain morphometry using structural MRI scans, Val/Met individuals have repeatedly been shown to have a smaller hippocampal volume relative to controls who are homozygous for Val allele (Val/ Val) (Pezawas et al. 2004; Szeszko et al. 2005; Bueller et al. 2006). This difference may be related to the role that BDNF and its receptors play in the development as well as continued plasticity of the brain (Huang and Reichardt 2001; Lu et al. 2005). In addition to hippocampus, studies have shown decreased volume in the dorsolateral prefrontal cortex, an area associated with planning and higher-order cognitive functioning, in carriers of the Met allele (Pezawas et al. 2004). It was also found that Met allele carriers have smaller temporal and occipital lobar gray matter volumes (Ho et al. 2006). Interestingly, the effect of  $BDNF_{Met}$  SNP on brain structure is modified by neuropsychiatric disorders. The bipolar disorder patients who carried the BDNF<sub>Met</sub> allele had the smallest hippocampus volumes compared to individuals without bipolar disorder and Val/Val homozygotes, and these decreases are most prominent in left anterior hippocampus (Chepenik et al. 2009; Matsuo et al. 2009). Within the bipolar cohort, individuals with one or more BDNF Met alleles showed greater losses in prefrontal gyrification index (GI) over time (Mirakhur et al. 2009). These studies suggest that the BDNF<sub>Met</sub> SNP may play a more prominent role in brain structure differences in subjects affected with bipolar disorder. Within schizophrenia patients, Met allele carriers had significantly greater reductions in frontal gray matter volume than Val-homozygous patients (Ho et al. 2007). Major depression patients had significantly smaller hippocampal volumes compared with controls and significantly smaller hippocampal volumes were observed for patients carrying the Met-BDNF allele compared with subjects homozygous for the Val-BDNF allele, which suggest that Met-BDNF allele carriers might be at risk to develop smaller hippocampal volumes and may be susceptible to major depression (Frodl et al. 2007).

Brain structure alterations associated with  $BDNF_{Met}$  SNP were further analyzed in  $BDNF_{Met}$  knock-in mouse. Using Cavalieri volume estimation, a significant decrease (about 14 %) in hippocampal volume in  $BDNF_{Met}$  mice was observed, which was comparable to the decrease in the humans carriers of Met allele (Pezawas et al. 2004; Chen et al. 2006). Ventromedial prefrontal cortex (vmPFC) volume was also found to be significantly decreased in  $BDNF_{Met/Met}$  mice compared with wild-type mice (Yu et al. 2009). Golgi staining and fractal dimension analyses further revealed a significant decrease in dendritic complexity in the dentate gyrus and vmPFC neurons from  $BDNF_{Met}$  mice (Chen et al. 2006; Yu et al. 2009), which might in part account for the decreased volume in  $BDNF_{Met}$  carriers.

# 5 Behavior Phenotypes Associated with BDNF<sub>Met</sub> Polymorphism

One of the common clinical symptoms among the psychiatric disease affected by the BDNF<sub>Met</sub> variant is the impairment of higher cognitive abilities. Individuals with the Val/Met genotype have been shown to be associated with poorer episodic memory in comparison with Val homozygotes, which rely heavily on the hippocampus (Egan et al. 2003; Hariri et al. 2003). Moreover, functional magnetic resonance imaging (fMRI) studies showed Met carriers exhibited relatively diminished hippocampal activation compared with Val/Val individuals during both the encoding and retrieval processes of a declarative memory task (Egan et al. 2003; Hariri et al. 2003). These findings taken together suggest that carriers of the Met allele have a selective impairment in hippocampal-dependent memory. In bipolar patients, the performance in Wisconsin card sort task (WCST) was significantly better in subjects with Val/Val-BDNF genotype compared with Val/Met genotype, which suggests a role of BDNF<sub>Met</sub> SNP in prefrontal cognitive function in bipolar illness (Rybakowski et al. 2003). In both patients with schizophrenia and healthy volunteers, Met allele carriers had poorer verbal memory performance than their Val-homozygous counterparts. In visuospatial abilities, Met allele-associated visuospatial impairment was specific to patients with schizophrenia but not healthy volunteers (Ho et al. 2006). These studies suggest that the association between the BDNF<sub>Met</sub> variant and poor medial temporal lobe-related memory performance and that the BDNF<sub>Met</sub> variant may have a specific role in conferring visuospatial dysfunction in schizophrenia (Ho et al. 2006). However, BDNF<sub>Met</sub> variant did not appear to impact general intellectual abilities or affect attention, problem solving, and language skills (Ho et al. 2006). A recent study also showed that BDNF<sub>Met</sub> polymorphism is associated with differences in brain motor system function, altered short-term plasticity, and greater error in short-term motor learning (McHughen et al. 2010).

The BDNF<sub>Met</sub> knock-in mouse provided a unique tool to assess the behavior phenotypes associated with BDNF<sub>Met</sub> polymorphism in a controlled genetic and environmental background which was not feasible in humans. In the fear conditioning test, BDNF<sub>Met</sub> mice showed significant less context-dependent memory than wild-type mice (Chen et al. 2006), which was consistent with the hippocampal-dependent learning defect in human carriers of the BDNF<sub>Met</sub> allele. When placed in conflict settings, BDNF<sub>Met/Met</sub> mice display increased anxiety-related behaviors in three separate tests (elevated plus maze, open field, and novelty-induced hypophagia) and thus provide a link between BDNF<sub>Met</sub> SNP and anxiety (Chen et al. 2006). Interestingly, the elevated anxiety in BDNF<sub>Met/Met</sub> mice could not be rescued by a common serotonin reuptake inhibitor (SSRI) fluoxetine, which suggests that humans with the BDNF<sub>Met</sub> allele may not have optimal responses to this class of antidepressants. The BDNF<sub>Met/Met</sub> mouse may serve as a valuable model to identify novel pharmacologic approaches to treat anxiety symptoms that underlie many neuropsychiatric disorders.  $BDNF_{Met/Met}$  mouse was further found to exhibit abnormalities in aversive memory extinction (Yu et al. 2009). This abnormality in extinction learning may be explained by alterations in neuronal morphology, as well as decreased neural activity in the vmPFC. Recently, Soliman et al. reported the impaired extinction learning in human  $BDNF_{Met}$  allele carriers relative to non-Met allele carriers, which was paralleled by atypical frontoamygdala activity in humans (Soliman et al. 2010). Understanding the effect of BDNF<sub>Met</sub> allele on impaired learning of cues that signal safety versus threat provided insight into the role of BDNF<sub>Met</sub> SNP in anxiety disorders. In BDNF<sub>Met/Met</sub> mouse, D-cycloserine (DCS), a partial agonist of NMDA receptor, was effective in rescuing the extinction defect (Yu et al. 2009), suggesting that when coupled with behavior therapy, DCS may be an effective treatment option for anxiety disorders in humans with this genetic variant BDNF.

# 6 Conclusion

Brain-derived neurotrophic factor (BDNF) plays important roles in activity-dependent plasticity processes and has been implicated in the pathophysiology of affective disorders. The human genetic variant BDNF (Val66Met) represents the first example of neurotrophin family member and has been linked to psychiatric disorders. Combined molecular and cellular, human behavioral and imaging, and transgenic mouse approaches were employed to assess the impact of BDNF<sub>Met</sub> SNP on brain structure and function. The mouse model provides the opportunity to test dose-dependent effects of the BDNF<sub>Met</sub> allele in a controlled genetic and environmental background not feasible in humans. The human behavioral and imaging findings provide confidence that crossspecies translation is biologically valid. BDNF<sub>Met</sub> variant has been associated with cognitive impairment and anxiety; drug discovery strategies to increase BDNF release from synapses, prolong the half-life of secreted BDNF, or directly activate TrkB receptors may provide greater therapeutic responses for human populations with this common BDNF polymorphism. In addition, the BDNF<sub>Met/Met</sub> mouse may serve as a valuable model to evaluate novel pharmacologic strategy in humans with this polymorphic allele and may ultimately guide personalized medicine for related clinical disorders.

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# Blood Brain-Derived Neurotrophic Factor Levels and Mood Disorders

# Reiji Yoshimura and Jun Nakumura

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### Abstract

Blood levels of brain-derived neurotrophic factor (BDNF) plays an important role in mood disorders. It is considered that blood (serum or plasma) BDNF could be a candidate biological state marker for depressive state. In addition, blood BDNF levels might also reflect psychological job stress in healthy workers.

R. Yoshimura (🖂) • J. Nakumura

Department of Psychiatry, School of Medicine, University of Occupational and Environmental Health, Yahatanishi-ku, Kitakyushu, Japan e-mail: yoshi621@med.uoeh-u.ac.jp

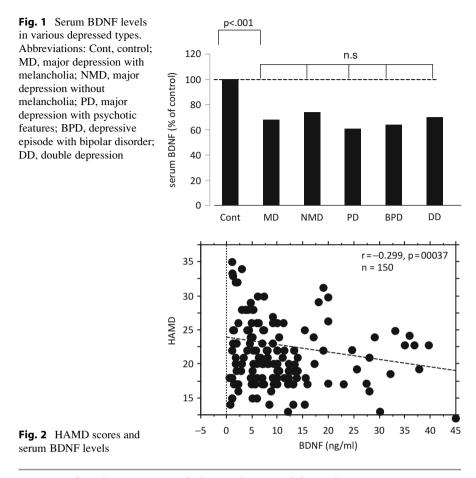
#### Keywords

Brain-derived neurotrophic factor • Serum • Plasma • Mood disorders • BDNF gene Val66Met

### 1 Introduction

Depression is heterogeneous and is diagnosed on the basis of a patient's symptoms, not on the basis of a laboratory test. Therefore, it is important to search for biological markers for depression due to the need of finding diagnostic adjuncts. Brain-derived neurotrophic factor (BDNF), a major neurotrophic factor, has been found to play a critical role in long-term potentiation, a cellular mechanism of learning and memory, suggesting that it can influence neuroplasticity (Fignov et al. 1996). BDNF is also needed for the survival and guidance of neurons during development as well as the survival and function of neurons during adulthood (Duman et al. 2000). The atrophy and loss of hippocampal or cerebral cortical neurons or of glia could result from a stress-induced loss of neurotrophic factors, from other processes that compromise neuronal function and activity, or from other insults depending on the patient's genetic background (Shelton 2000). There is growing evidence that BDNF might have a crucial role in depression (Duman et al. 2000; Mowla et al. 2001). The source of circulating BDNF remains unknown. Platelets, brain neurons, and vascular endothelial cells are currently considered to be putative sources. It was demonstrated that BDNF crosses the blood-brain barrier (Pan et al. 1998) and that BDNF levels in the brain and serum have been shown to undergo similar changes during the maturation and aging process in rats (Karege et al. 2002). Furthermore, Lang et al. reported that serum BDNF concentrations reflect some aspects of neuronal plasticity, as indicated by the association of BDNF levels with those of N-acetyl-aspartate, which reflects neuronal damage in the cerebral cortex (Lang et al. 2007). These results indicated that blood BDNF levels might in part reflect the BDNF levels in the brain. Moreover, Klein et al. (2011) recently reported that measures of blood and plasma BDNF levels reflect brain-tissue BDNF levels. We reported that serum BDNF levels are approximately 20 times greater than those in platelet-poor plasma, and a close correlation was found between plasma and serum levels (Yoshimura et al. 2010a). Platelets are the major source of serum BDNF, as they sequester large quantities of it for release during clotting. Sources of circulating plasma BDNF include vascular endothelial cells as well as brain cells (Lommatzsch et al. 2005). Platelets have a life span of approximately 10 days, whereas plasma BDNF is minimally affected by the amount stored in platelets, and therefore is likely to be a more suitable index of brain BDNF levels (Marano et al. 2007) (Fig. 1).

In this chapter, the role of serum BDNF and its association with mood disorders predominantly will be discussed mainly based on our previous findings.



## 2 Blood BDNF Levels in Patients with Major Depressive Disorder

The serum BDNF levels were significantly lower in those patients than in the healthy subjects: No difference in serum BDNF levels between with melancholia and without melancholia. In addition, no difference was observed in serum BDNF levels between the patients with psychotic features (psychotic depression) and without ones. There have been at least three meta-analyses of studies of blood BDNF levels in patients with major depressive disorder (Bocchio-Chiavetto et al. 2010; Brunori et al. 2010; Sen et al. 2008). The results of those investigations have consistently revealed that blood BDNF levels in patients with major depressive disorder are significantly lower than those of healthy subjects. A negative correlation was also found between the scores of Hamilton Rating Scale for Depression (HAMD) and serum BDNF levels, which means serum BDNF levels reflect the severity of depressive state (Yoshimura et al. 2011; Shimizu et al. 2003). From these findings, it is deduced blood BDNF level is a candidate biomarker for depression (Fig. 2).

### 3 Blood BDNF Levels in Patients with Bipolar I Disorder

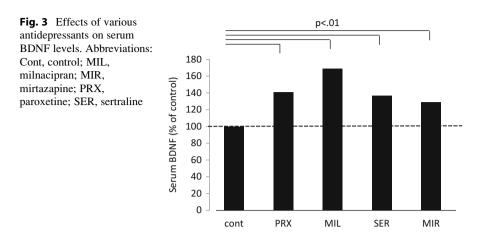
Plasma BDNF levels in depressive episodes were significantly lower than those in manic episodes in patients with bipolar I disorder or healthy subjects (Yoshimura et al. 2006). Fernandes et al. recently performed meta-analysis including our study, and they concluded that blood BDNF levels are consistently reduced during maniac and depressive episodes, but not during euthymic state in patients with bipolar I disorder (Fernandes et al. 2011).

## 4 Blood BDNF Levels in Patients with Dysthymic Disorder

Dysthymic disorder has a number of typical characteristics such as low drive, low self-esteem, and a low capacity for pleasure. They will usually find little pleasure in usual activities and pastimes (Griffiths et al. 2000). In addition, BDNF levels among patients with dysthymic disorder, major depressive disorder, and healthy controls were measured. Serum BDNF levels in the dysthymic disorder group were significantly lower than those in the healthy controls. Moreover, no difference was found between the dysthymic disorder group and the major depressive disorder group (Yoshimura et al. 2011). On the other hand, Aydemir et al. (2007) reported that serum BDNF concentrations of the dysthymic group were significantly higher than that of the major depressive disorder group, and it was not different from the level of the control group. One of the reasons of the discrepancy of the results might be due to the difficulty of the diagnosis of dysthymic disorder. Actually, it is difficult to diagnose patients with dysthymic disorder because of the subtle nature of the symptoms and the patients often can hide them in social situations, making it challenging for others to detect symptoms. Additionally, dysthymic disorder often occurs at the same time as other psychological disorders which adds a level of complexity to determine the presence of dysthymic disorder, particularly because there is often an overlap in the symptoms of other psychiatric disorders (Sansone et al. 2009).

# 5 Blood BDNF Levels in Patients with Major Depressive Disorder with Alcohol Dependence

Alcohol dependence is often comorbid with major depressive disorder (Regiet et al. 1990). Alcohol use in depression patients increases their severity of depressive state and suicidal risk. Zanardini et al. (2011) reported that serum BDNF levels were decreased in patients with alcohol dependence However, no reports have been found comparing blood BDNF levels with and without alcohol dependence in depressed patients. Serum BDNF levels in the patients with or without alcohol dependence were significantly lower than those in the healthy controls. There was however no difference in serum BDNF levels between the depressed patients with and without alcohol dependence (Umene-Nakano et al. 2009).



# 6 Effects of Antidepressants on Serum BDNF Levels in Depressive Patients

Recent review by Molendijk et al. demonstrated that St John's wart and selective serotonin reuptake inhibitor (SSRI), but not serotonin noradrenaline reuptake inhibitor (SNRI), tricyclic antidepressant (TCA), or noradrenergic and specific serotonergic antidepressants (NaSSAs) increased serum BDNF levels (Molendijk et al. 2010). Effects of paroxetine, a SSRI, and milnacipran, a SNRI, on serum BDNF levels in patients with major depressive disorder were investigated. Treatment with paroxetine and milnacipran for at least 8 weeks, but not 4 weeks, both increased serum BDNF levels in responders, but not in nonresponders to the same degree (Yoshimura et al. 2007). Taken together, serotonin rather than noradrenaline might play an essential role for increasing BDNF (Molendijk et al. 2010) (Fig. 3).

# 7 Effects of Repetitive Transcranial Magnetic Stimulation (rTMS) or Electrical Convulsive Therapy (ECT) on Blood BDNF Levels

rTMS or ECT is effective for treatment-resistant depression. A systematic review by Lam et al. (2008) reported that the pooled response and remission rate rates were 25 % and 17 %, respectively. ECT is widely used for treatment in major depressive disorder, bipolar disorder, and schizophrenia. The mechanisms of rTMS and ECT have not been fully elucidated. To the best of our knowledge, this is the first report that rTMS increased plasma BDNF levels at least 5 weeks in responders with treatment-resistant depressed patients (Yukimasa et al. 2006). Furthermore, ECT also increased serum BDNF levels in responders with refractory patients within 4 weeks (Okamoto et al. 2008). Taking these findings into account, it can be deduced that rTMS or ECT rapidly and robustly might increase blood BDNF levels

in depressed patients. Bocchio-Chiavetto et al. have demonstrated that ECT significantly increased serum BDNF levels in paralleled with clinical improvement of depressive symptoms (Bocchio-Chiavetto et al. 2006). On the other hand, Fernandes et al. (2009) reported that no changes were observed in the serum BDNF before and after the ECT treatment. Zanardini et al. (2006) reported that a significant increase of serum BDNF was found after rTMS treatment. It still remains controversial: How rTMS or ECT increased blood BDNF levels in depressed patients. Farther replications in larger sample will be performed.

# 8 Polymorphism of BDNF Gene Val66Met and Response to SSRIs

The relationship between the BDNF gene polymorphism (Val66Met) and the clinical response of patients with major depressive disorder to selective serotonin reuptake inhibitors (SSRIs; paroxetine and sertraline) was investigated. In addition, serum BDNF levels in these patients were considered together with the clinical response. A negative association was found between the serum BDNF levels and the Hamilton Rating Scale for Depression (HAMD) scores. No correlation was observed between BDNF gene Val66Met polymorphism and response to SSRIs or between BDNF Val66Met polymorphism and serum BDNF levels. These results suggest that serum BDNF level is a state biomarker for depression, but that in depressed patients, presence of the BDNF Val66Met polymorphism is independent of both the response to SSRI treatment and serum BDNF levels (Yoshimura et al. 2011). In contrast, recent meta-analysis by Zou et al. (2010) demonstrated that the association between BDNF Val66Met polymorphism and treatment response in patients with major depressive disorder, and Val66Met heterozygous patients has a better response rate in comparison to Val66Val homozygote patients, especially in the Asian population.

## 9 Adding an Atypical Antipsychotic Drug to an Antidepressant Increased Rapidly Plasma BDNF Levels in Treatment-Resistant Depressed Patients: Preliminary Findings

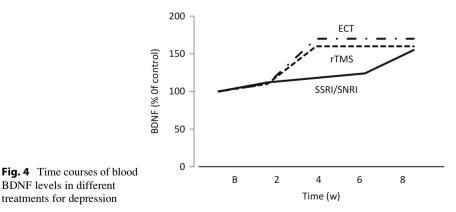
Only two-thirds of depressive patients respond to antidepressant treatment. Addition of an atypical antipsychotic drug to ongoing treatment with an antidepressant has been considered effective and well-tolerated, and that the improvements of depressive symptoms were related with the increase of plasma BDNF levels after adding the atypical antipsychotic drug to the antidepressant (Yoshimura et al. 2010b). No relationship was however observed between BDNF gene Val66Met polymorphism and the response to the atypical antipsychotic drug to the antidepressant. To the best of our knowledge, this is the first report demonstrating an atypical antipsychotic drug to an antidepressant enhances plasma BDNF levels in refractory depressive patients.

## 10 Longitudinal Follow-Up Study of Serum BDNF Levels and the Relapse of Depressive Episodes

Long-term study of serum BDNF levels in patients with major depressive disorder. A longitudinal follow-up study measuring serum BDNF levels after remission of depressive episodes. Serum BDNF levels at relapse were significantly lower than those at remission or after 6 months after remission. No difference was observed between levels at remission and at 6 months after remission. These results indicate that serum BDNF levels at 6 months after remission might not be associated with subsequent depressive episodes.

# 11 No Association Between Responses to the Addition of an Atypical Antipsychotic Drug to a SSRI or SNRI and BDNF (Val66Met) Polymorphism in Refractory Major Depressive Disorder: Preliminary Study

The association between the BDNF (Val66Met) polymorphism and the response to the addition of an atypical antipsychotic drug to a SSRI or SNRI in treatment-refractory depression was observed. Forty-two patients meeting the DSM-IV criteria for major depressive disorder and who were treated with at least two courses of a single antide-pressant, but whose HAMD-17 scores were 15 points or more and were reduced less than 50 % over at least a 4-week treatment periods, were enrolled in the study. No correlation was found between the BDNF (Val66Met) polymorphism and a positive response to the atypical antipsychotic addition. In addition, patients who experienced remission were followed up 1 year after discontinuation of the atypical antipsychotic drug at least within 3 months. Eleven of 19 patients (47 %) relapsed within a year. These results suggest that the BDNF (Val66Met) polymorphism is not associated with the response to the augmentation of a SSRI or SNRI with an atypical antipsychotic drug, and the combination regimen of an atypical antipsychotic drug and a SSRI or SNRI should be continued for 3 months or more in refractory depressed patients in the Japanese population (Fig. 4).



## 12 Mental Stress, Personality Trait, and Blood BDNF Levels

Lang et al. first reported a negative correlation between the BDNF serum concentration and the depression-related factor neuroticism (Yoshimura et al. 2011). These results indicate that low BDNF levels in healthy humans with depressive personality traits might constitute a risk marker, reflecting a personality profile that is linked to vulnerability to mood disorders. Recently, Terracciano et al. also reported that serum BDNF might represent a biological correlate of neuroticism and not just of transient depressive state (Lang et al. 2004). Associations between plasma BDNF levels and personality trait evaluated by using NEO-FFI (neuroticism, extroversion, openness, agreeableness, and conscientiousness) also were examined. A positive correlation was observed between plasma BDNF levels and extroversion, but not neuroticism (Terracciano et al. 2011). The discrepancy of our results and those in other two studies remain controversial. One of the reasons might be the difference in the ethnicity. An association between psychological job stress and serum BDNF levels in Japanese hospital employee also was examined. Psychological job stress was evaluated. A negative correlation was observed between the scores of psychological stress and serum BDNF levels (Mitoma et al. 2008; Okuno et al. 2011). In other words, subjects with high psychological job stress had low BDNF levels. These results suggest that serum BDNF levels reflect psychological job stress in healthy workers.

## 13 Conclusion

Blood (serum and plasma) BDNF levels are useful state biomarker for depressive stateon. Blood BDNF levels reflect the severity of depressive state. SSRIs and SNRIs increased serum BDNF levels in the responders with major depressive disorder. Adding an atypical antipsychotic drug to a SSRI or a SNRI improved clinical symptoms in patients with refractory depression, which was accompanied with the increase plasma BDNF levels. The polymorphism of BDNF gene Val66Met was neither associated with the response to SSRIs nor an atypical augmentation to SSRIs or SNRIs. Further research should be done to reconfirm our preliminary findings.

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# Neuroprotection in Demyelinating Diseases: The Therapeutic Potential of the Neurotrophins

# Junhua Xiao, Richard A. Hughes, and Simon S. Murray

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#### Abstract

The myelin sheath has evolved to exert critical influences upon the central and peripheral nervous systems. Recently, the neurotrophins have been implicated in influencing the dynamic and complex signals that occur between neurons and myelinating glial cells – Schwann cells in the peripheral nervous system and oligodendrocytes in the central nervous system – that regulate myelination. Somewhat surprisingly, the neurotrophins have been found to influence

J. Xiao (🖂) • S.S. Murray

Department of Anatomy and Neuroscience, The University of Melbourne, Parkville, VIC, Australia e-mail: xiaoj@unimelb.edu.au

R.A. Hughes Department of Pharmacology, University of Melbourne, VIC, Australia

myelination in a complex manner involving both promyelinating and inhibitory signals that can be directed against either neuronal or glial cells that ultimately regulate central and peripheral myelin formation in distinct ways. The neurotrophins and their receptors have also been shown to influence the severity and affect remyelination in distinct in vivo models of demyelinating disease. Together, these data indicate that the selective targeting of neurotrophin receptors to promote remyelination offers an exciting prospect for the treatment of demyelinating diseases.

List of Abbreviations					
CGRP	Calcitonin gene-related peptide				
CNS	Central nervous system				
DPN	Diabetic peripheral neuropathy				
DRG	Dorsal root ganglion				
EAE	Experimental autoimmune encephalomyelitis				
EAN	Experimental autoimmune neuritis				
Erk	Extracellular signal-regulated kinase				
GPCR	G Protein-coupled receptor				
MAPK	Mitogen-activated protein kinase				
MS	Multiple sclerosis				
NRG	Neuregulin				
NTR	Neurotrophin receptor				
OPCs	Oligodendrocyte progenitor cells				
PNS	Peripheral nervous system				
TrkB	Tropomyosin-related kinase B				

# 1 Introduction

The myelin sheath is critical for normal nervous system function. Its evolution has effectively given the nervous system enhanced compactness, reduced its metabolic demands, improved action potential synchrony, and allowed the rapid processing of complex information. This has been achieved via at least two distinct and critical influences on the nervous system. The first is to increase the speed at which the action potential is propagated along the axon, by permitting saltatory conduction. Electrophysiological studies have clearly shown the impact this has on nervous system function, demonstrating that nervous system demyelination results in slowing of the conduction velocity of action potentials and in more severe cases can actually lead to conduction block (McDonald and Sears 1969). While these studies theoretically accounted for the neurological deficits encountered in people suffering from demyelinating diseases, the clinical picture appeared more complex. Brain imaging studies revealed that white matter lesion load did not necessarily correlate with the extent of neurological deficit. Indeed,

histological studies have now revealed that injured and transected axons are present within both demyelinated and actively demyelinating lesions and that progressive axonal loss resulting from demyelination is a key contributor to clinical decline (Trapp et al. 1997; Bitsch et al. 2000). It is now increasingly apparent that the second critical influence that myelinating glial cells exert is to confer protective and trophic support to the axons which they ensheathe (Yin et al. 2006). Collectively, these studies indicate that there is an intimate relationship between damage to either the myelinating glial cell or the myelin sheath itself and axonal degeneration. The evolution of the myelin sheath and saltatory conduction reduced the metabolic demands and energy expenditure required by the nerve cell to relay the action potential down its axon, and perhaps this at least in part contributes to the axonal-protective properties of myelin. However it is also clear that the myelinating cells support axonal integrity and survival independent of the myelin sheath itself (Lappe-Siefke et al. 2003; Lee et al. 2012; Oluich et al. 2012). Thus, the protective influences that myelinating glial cells and the myelin sheath exert on axonal integrity suggest that any successful therapy against demyelinating diseases must consider both survival of myelinating glial cells and myelin regeneration.

In human demyelinating diseases such as multiple sclerosis (MS) and some peripheral demyelinating neuropathies, the myelinating glial cells progressively die, disrupting these myelin sheaths and placing axonal integrity at risk. The body has an innate mechanism for remyelination, whereby myelin sheaths are restored to the damaged axons, protecting them from further degeneration (Irvine and Blakemore 2008; Franklin et al. 2012). In MS, this is affected by regenerating oligodendrocytes through the generation, recruitment, and differentiation of OPCs resident within the CNS. However, remyelination is often incomplete, becomes less efficient with age, and following successive demyelinating events is ultimately insufficient and invariably leads to irreversible axonal damage and progressive and extensive disability (De Stefano et al. 1998).

In these diseases an autoimmune response is commonly observed, which is generally believed to contribute to the extent of demyelination. While immunetargeted therapies are available to limit demyelination, no treatments are currently available to directly promote myelin repair. Intuitively, studies that examine the growth factors and molecular mechanisms that promote myelination during development and remyelination following injury are essential if the therapeutic targets that promote myelin repair and thereby limit axonal injury and ultimately disability are to be identified. Recent reviews have examined the neurotrophin family of growth factors and the roles they play in regulating myelination during development in detail (Rosenberg et al. 2006; Xiao et al. 2009a). We briefly review here the current understanding of the mechanisms that the neurotrophins and their receptors utilize to influence myelination during development and identify the protective roles they exert in demyelinating diseases. Finally we discuss the evolving potential for the selective targeting of neurotrophin receptors for therapeutic benefit in the context of myelin repair following a demyelinating insult.

## 2 The Neurotrophins, Their Receptors, and Myelination During Development

### 2.1 The Neurotrophins and Their Receptors

The neurotrophin family of growth factors comprises nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), neurotrophin-3 (NT-3), and neurotrophin-4/5 (NT-4/5). They are widely expressed and exist in two distinct forms. They are initially synthesized as larger precursor proteins, known as proneurotrophins, which subsequently undergo proteolytic cleavage either intracellularly or extracellularly to yield the mature neurotrophins. It is generally accepted that the proneurotrophins modulate cell death (Teng et al. 2010); however, there is emerging evidence they can also exert distinct influences including modulating synaptic plasticity and axonal competition at developing neuromuscular synapses (Woo et al. 2005; Je et al. 2012). To date, there is no evidence to indicate that the proneurotrophins directly influence myelination, whether central or peripheral.

In contrast, it is well established that the mature neurotrophins exert profound and diverse effects upon neural development as well as in maintaining and modulating the adult mature nervous system. The mature neurotrophins signal through two distinct classes of structurally unrelated transmembrane receptors: the receptor tyrosine kinase tropomyosin-related kinase (Trk) family and p75NTR (Chao 2003). These receptors exhibit a distinct capacity to interact with the neurotrophins. Whereas all the neurotrophins can bind nonselectively and with approximately the same affinity to p75NTR, they display selectivity in their interaction with Trk receptors: NGF binds selectively to TrkA, BDNF and NT4/5 to TrkB, and NT-3 to TrkC (Chao 2003; Huang and Reichardt 2003). Increasing the complexity of these ligand-receptor interactions, when Trk and p75NTR are co-expressed on the same cell, they can also form a receptor complex wherein p75NTR modulates the affinity of Trk receptor binding and selectivity of Trk receptor activation (Esposito et al. 2001). Despite this complexity, genetic analysis of mice deficient in either the neurotrophins or their receptors reveals that, by and large, Trk receptor signalling transduces the key influences of the neurotrophins (Conover and Yancopoulos 1997; Huang and Reichardt 2003).

#### 2.2 The Neurotrophins Regulate Schwann Cell Myelination

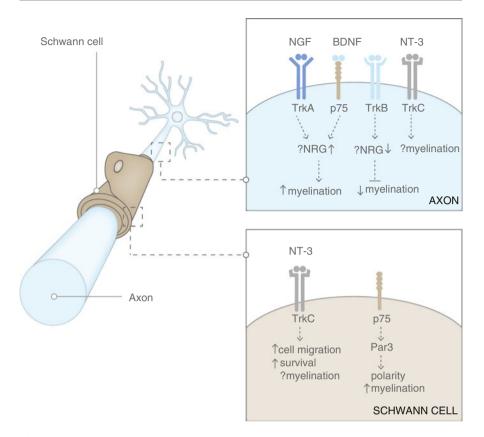
It is well established that both peripheral neurons and Schwann cells express a variety of neurotrophin receptors, as well as expressing and secreting neurotrophins. Analyses of either neurotrophin or neurotrophin receptor knockout mice reveal the absolute requirement of the ligands and their receptors for peripheral neuron development, showing that  $\sim 80$  % of DRG neurons are dependent on NGF for survival (Crowley et al. 1994),  $\sim 30$  % are dependent on BDNF (Jones et al. 1994), and  $\sim 70$  % are dependent on NT-3 (Liebl et al. 1997; Tessarollo et al. 1997).

Consequently, while it is clear that the neurotrophins and their receptors exert critical influences on the development of the peripheral nervous system, analysis of the precise effects that these molecules have on peripheral myelination in vivo has been problematic. However, development and manipulation of in vitro myelination assays has provided insight into the roles the neurotrophins and their receptors are playing in this context. Somewhat surprisingly, despite Schwann cells expressing several neurotrophin receptors, the key influences that the neurotrophins exert on peripheral myelination appear to be directed through neuronally expressed receptors that ultimately regulate axonal receptivity to myelination by Schwann cells.

Chan and colleagues were the first to identify that NGF exerts a positive influence upon Schwann cell myelination. Through the use of in vitro myelination assays, they identified that NGF enhanced Schwann cell myelination specifically of TrkA-expressing DRG neurons. This influence of NGF was specific to TrkA, as NGF also promoted myelination of neurons not expressing p75NTR and exerted no influence on TrkB-expressing DRG neurons (where TrkA expression is absent) (Chan et al. 2004). Thus, NGF exerts its promyelinating influence selectively through neuronally expressed TrkA receptors (Fig. 1). Contextually this fits well developmentally, as  $\sim 80 \%$  of DRG neurons express TrkA during the early postnatal period when peripheral nerves are actively being myelinated. Thus, TrkA is well placed to direct key influences upon DRG neurons to regulate their myelination in vivo (Wiggins et al. 1975; Silos-Santiago et al. 1995; Molliver and Snider 1997; Xiao et al. 2009a). Interestingly, TrkA expression is strikingly developmentally regulated, in that while  $\sim 80$  % of DRG neurons express TrkA during embryogenesis and early postnatal life, the expression of TrkA is remarkably downregulated in adult DRG neurons (Molliver and Snider 1997). This suggests that the utility of NGF as a potential remyelination therapy in adults may be limited.

The influence that BDNF exerts upon peripheral myelination is considerably more complex, with evidence demonstrating that BDNF can promote or inhibit the myelination of distinct subsets of DRG neurons. Like NGF, BDNF regulates Schwann cell myelination through activation of neuronally expressed receptors. However, the precise influence that BDNF exerts is dependent on the subset of neurotrophin receptors expressed on the particular DRG neuron. BDNF inhibits Schwann cell myelination of TrkB+ DRG neurons; however, in the absence of neuronal TrkB, BDNF can promote the myelination via activation of neuronal p75NTR (Chan et al. 2001; Cosgaya et al. 2002; Xiao et al. 2009b). The contrasting influence that BDNF exerts upon Schwann cell myelination is therefore context specific, due to BDNF activating distinct neuronal receptors – p75NTR to promote myelination and TrkB to inhibit it (Xiao et al. 2009b) (Fig. 1). For this reason, the utility of BDNF itself as a potential remyelination therapy in adults may also be limited. This also reveals a fundamental difference as to how NGF and BDNF promote Schwann cell myelination: NGF promotes myelination via activation of neuronal TrkA receptors (Chan et al. 2004), whereas BDNF promotes myelination via activation of neuronal p75NTR receptors (Xiao et al. 2009b) (Fig. 1).

In contrast to the situation for NGF and BDNF, there have been relatively few studies that have attempted to identify the precise influence that NT-3 exerts on



**Fig. 1** Neurotrophin influences on Schwann cell myelination. Top panel: Neuronal influences of the neurotrophins. NGF exerts a promyelinating influence via TrkA activation. BDNF promotes myelination via p75NTR activation, but inhibits myelination via TrkB. NGF and BDNF putatively differentially regulate neuregulin (*NRG*) expression to regulate myelination. The precise influence that neuronal TrkC receptors exert is unclear. *Bottom panel*: Neurotrophin influences on Schwann cells. NT-3 effects Schwann cell migration and survival, whereas BDNF influences Schwann cell polarity in a p75NTR-dependent manner. The precise role these influences exert on myelination is unclear

Schwann cell myelination. In vitro, exogenous NT-3 inhibits myelin formation, and injection of NT-3 adjacent to the sciatic nerve of neonatal mouse pups also results in reduced myelin protein formation, supporting an inhibitory role for NT-3 upon peripheral myelination, most likely via activation of the TrkC receptor (Chan et al. 2001; Cosgaya et al. 2002) (Fig. 1). However, as Schwann cells and some DRG neurons express TrkC, the precise mechanism that NT-3 utilizes to influence myelination remains unclear.

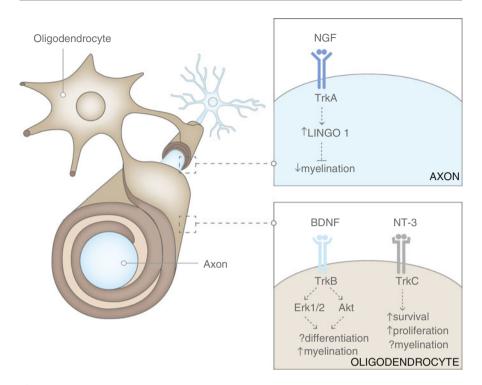
Collectively, these data demonstrate that the influences the neurotrophins exert upon the myelination of peripheral axons are highly dependent on the complement of neurotrophin receptors expressed at critical times in development. It is clear that at least NGF and BDNF direct neuronal events that alter axonal receptivity to Schwann cell myelination. Importantly, the critical neuronal event that determines the ensheathment fate of peripheral axons is the expression of neuregulin-1 type III (Taveggia et al. 2005; Nave and Salzer 2006; Birchmeier and Nave 2008). Thus, it is likely that neurotrophin signalling within the distinct subsets of DRG neurons differentially regulates the expression of neuregulin-1 type III. However, this has yet to be definitively demonstrated.

It is intriguing that Schwann cells also express several neurotrophin receptors, including p75NTR, TrkC, and truncated TrkB receptors. Emerging data identify that the neurotrophins also exert direct influences on Schwann cells. BDNF can regulate Schwann cell migration (Yamauchi et al. 2004) as well as Schwann cell polarity and alignment to axons in a p75NTR-dependent manner (Chan et al. 2006) (Fig. 1). NT-3, acting through Schwann cell-expressed TrkC receptors, also promotes Schwann cell migration (Yamauchi et al. 2003; Yamauchi et al. 2005a, b), and analysis of NT-3 mutant mice suggests it also promotes survival of maturing Schwann cells (Woolley et al. 2008) (Fig. 1). These events all ultimately impact upon the process of myelination and clearly suggest that neurotrophins exert distinct influences directly upon both neurons and Schwann cells that collectively contribute to the orchestration of peripheral myelination.

#### 2.3 The Neurotrophins Regulate Oligodendrocyte Myelination

As in the peripheral nervous system, a variety of neurotrophins and neurotrophin receptors are expressed by both the neurons and oligodendrocytes of the central nervous system. Analyses of central nervous system myelination in either neurotrophin or neurotrophin receptor knockout mice have also proven largely problematic, as many of these mice die in the embryonic or early postnatal period prior to CNS myelination being complete. In any case, analyses of these mice are confounded by the global deletion, whereupon the potential selective influences that neuronal and oligodendroglial receptors exert upon myelination cannot be discriminated. In vitro myelination assays have again proved critical here, providing insight into the influences that the neurotrophins and their receptors exert upon oligodendrocyte myelination. Perhaps surprisingly, the data here clearly indicate that the neurotrophins exert quite distinct influences upon peripheral and central nervous system myelination.

The first indication that the neurotrophins did not exert uniform influences upon peripheral and central myelination came with the study of NGF. Chan and colleagues, adopting the same approach that demonstrated that NGF promoted Schwann cell myelination, found that NGF inhibited oligodendrocyte myelination (Chan et al. 2004). Mechanistically NGF was shown to be acting via the activation of neuronally expressed TrkA receptors in both cases but with starkly contrasting outcomes (Fig. 2, top panel). This reaffirmed what had been known for some time – that axonal control over peripheral and central nervous system myelination is mediated via unique and mutually exclusive mechanisms – but surprisingly



**Fig. 2** Neurotrophin influences on oligodendrocyte myelination. Top panel: Neuronal influences of the neurotrophins. NGF inhibits oligodendrocyte myelination via regulating LINGO-1 expression. *Bottom panel*: Neurotrophin influences on oligodendrocytes. BDNF activates TrkB to promote oligodendrocyte myelination, potentially via Erk1/Erk2 or Akt activation. NT-3 effects oligodendrocyte survival and proliferation; however, the precise influence it exerts on myelination is unclear

revealed that a common growth factor could exert opposite effects upon myelination. In this context, NGF regulates axonal expression of LINGO-1, a membrane-bound protein that acts as a negative regulator of oligodendrocyte but not Schwann cell myelination (Mi et al. 2005; Lee et al. 2007) (Fig. 2, top panel). As an aside, LINGO-1 exerts profoundly negative influences on oligodendrocyte myelination (Mi et al. 2005), and LINGO-1 antagonists have been found to promote remyelination in animal models of CNS demyelination (Pepinsky et al. 2011a, b). A phase 1 clinical trial utilizing a LINGO-1 antagonist has recently been completed.

Analysis of the impact that BDNF exerted on oligodendrocyte myelination also revealed marked differences to its influence on Schwann cell myelination. Both BDNF knockout and heterozygous mice display a hypomyelinated phenotype in the CNS (Cellerino et al. 1997; Vondran et al. 2010; Xiao et al. 2010), suggesting a key role for BDNF in promoting oligodendrocyte myelination. This was verified in vitro, with BDNF found to promote myelination via a direct effect upon oligodendrocyte-expressed TrkB receptors (Cohen et al. 1996; Du et al. 2003; Xiao et al. 2010) (Fig. 2). This was the first demonstration that a neurotrophin exerted its promyelinating influence by directly activating receptors expressed by the myelinating glial cell. This was subsequently verified in vivo through the analysis of mice in which TrkB had been deleted specifically in oligodendrocytes. This deletion significantly disrupted normal myelination in the CNS, resulting in a significant reduction in myelin protein expression and myelination of CNS white matter tracts during development (Wong et al. 2013). Interestingly, analyses revealed significantly reduced myelin thickness without oligodendrocyte number being affected, indicating that TrkB exerts an important influence in selectively promoting myelin ensheathment. These conditional knockout mice also exhibited an increased density of oligodendrocyte progenitor cells (OPCs) in CNS white matter tracts, which in vitro analyses suggested was dependent upon TrkC and p75 expression (Wong et al. 2013). This is another exemplar demonstrating the complexity of the neurotrophin signalling.

As the signalling by the Trk receptors is reasonably well characterized (Huang and Reichardt 2001), this had the potential to provide novel insight into the molecular control of myelination. Indeed, several canonical signalling pathways downstream of TrkB have been implicated in promoting oligodendrocyte myelination. Activation of both the MAPK (Chew et al. 2010; Fyffe-Maricich et al. 2011; Ishii et al. 2012) and Akt (Narayanan et al. 2009; Tyler et al. 2009) signalling pathways promote oligodendrocyte differentiation and myelination. Through screening candidate signalling pathways in in vitro myelination assays, it was found that BDNF-induced activation of extracellular signal-regulated kinase 1 and 2 (Erk1/Erk2) signalling in oligodendrocytes positively correlated with myelination (Xiao et al. 2012). Indeed, Erk1/Erk2 has been identified as a key signalling pathway that regulates the promyelinating effect of BDNF (Xiao et al., 2012; Ishii et al., 2012).

NT-3 is well known as an oligodendrocyte survival factor and was in fact one of the first growth factors identified to support the survival and proliferation of oligodendrocyte progenitor cells in vitro and in vivo (Barres et al. 1993, 1994; Kumar and de Vellis 1996; Kumar et al. 1998; Saini et al. 2004) (Fig. 2). It is important to note though that in addition to these effects upon progenitor cell survival, exogenous NT-3 also promotes oligodendrocyte myelination in vitro (Yan and Wood 2000; Rubio et al. 2004). The phenotype of the NT-3 knockout mouse supports a promyelinating role for NT-3 in vivo (Kahn et al. 1999). However, insufficient attention has been paid to NT-3 in this context, and whether its primary influence in the central nervous system in vivo is the generation and proliferation of OPCs or as a distinct promyelinating factor or both remains to be definitively established.

Similar to that observed for Schwann cell myelination, the influences the neurotrophins exert upon oligodendrocyte myelination are highly dependent on the complement of neurotrophin receptors expressed by the different cell types. While NGF initiates neuronal events that alter axonal receptivity to oligodendrocyte myelination, BDNF and NT-3 directly target oligodendrocyte-expressed

receptors to influence myelination. Somewhat surprisingly, despite p75NTR being expressed by both neurons and oligodendrocytes, it does not appear to exert a significant influence upon central nervous system myelination (Xiao et al. 2010), unlike that observed with Schwann cell myelination (Cosgaya et al. 2002; Chan et al. 2006; Xiao et al. 2009b).

Collectively, identification of the key influences that the neurotrophins exert on both peripheral and central nervous system myelination has revealed that their actions are highly contextual, with no unified response apparent. This is at least in part reflected in the complex nature with which the neurotrophins interact with their respective receptors but also reflects the distinct mechanisms that have evolved to generate central and peripheral myelin. From a therapeutic perspective this is potentially problematic, demanding that any potential neurotrophin-based approach be selective for central or peripheral remyelination, with the caveat that what might promote remyelination in the CNS could inhibit remyelination in the PNS. Nevertheless, neurotrophin-based approaches have been adopted to investigate their efficacy in promoting remyelination following a demyelinating injury to both the CNS and PNS.

# 3 The Neurotrophins, Their Receptors, and Remyelination Following Injury

Demyelination is the pathological process in which myelin sheaths are lost from around axons (Franklin and Ffrench-Constant 2008), with MS and Guillain-Barre syndrome (GBS) being the most common demyelinating diseases of the central and peripheral nervous systems, respectively (Miller 1996). Demyelination can result from a direct insult targeted at the myelin forming glial cells, which commonly occurs in CNS demyelination (Franklin and Ffrench-Constant 2008). The process of remyelination, in which entire myelin sheaths are restored to demyelinated axons, can reinstate saltatory conduction, resolve functional deficits, and promote axonal integrity and neuronal survival (Jeffery and Blakemore 1997; Liebetanz and Merkler 2006; Zawadzka and Franklin 2007). In recent years, multiple studies have examined the influence that the neurotrophins exert upon models of CNS autoimmune demyelination and remyelination. In contrast, there have been relatively few studies examining the influence that the neurotrophins exert upon models of peripheral nervous demyelination.

## 3.1 The Neurotrophins Promote Peripheral Nervous System Remyelination

Peripheral neuropathy encompasses complex neurological conditions, covering many disorders of the peripheral nervous system (Hughes 2008). The causes of peripheral neuropathy are protean: they can be either genetic in origin or acquired, with acquired neuropathies having either metabolic, ischemic, inflammatory,

or toxic precipitants (Hughes and Cornblath 2005; Hughes 2008). Peripheral neuropathies are also usefully classified as either axonal or demyelinative in origin. In axonal neuropathies the primary insult is directed to the neuron, and demyelination follows as a secondary consequence, whereas in demyelinative neuropathies the insult is directed to the Schwann cell that ensheathes the axon with myelin. Neuropathies are nevertheless not always pure; for example, there are circumstances in which the pathogenic insult is directed to both the axon and the Schwann cell; furthermore, in severe demyelinative neuropathy there is often secondary axonal degeneration because of the loss of trophic support that the Schwann cells otherwise provide.

Axonal peripheral neuropathy is most commonly caused by diabetes but can also result from a range of metabolic disorders, infectious agents, vasculitis, toxins, autoimmune and paraproteinemia neuropathies, and drugs. inherited polyneuropathies (McHugh and McHugh 2004). In experimental models of peripheral polyneuropathy, a reduction in the availability of the neurotrophins and alterations in the expression of their receptors has been observed, affecting both neuronal survival and maintenance of a myelinated phenotype (Apfel 1999a; Pittenger and Vinik 2003). The most widely utilized model of peripheral neuropathy is diabetic peripheral neuropathy (DPN) (Apfel et al. 1994, 1998; Apfel 1999a, b; Fressinaud et al. 2003; Pittenger and Vinik 2003). While DPN itself represents a heterogeneous variety of disorders affecting the peripheral nervous system, it is modelled by inducing diabetes in rats by administration of streptozocin.

DPN is characteristically described as degeneration of both small- and largediameter sensory fibers, axonal degeneration, demyelination and atrophy, and ultimately associated with failed axonal regeneration, remyelination, and synaptogenesis (Pittenger and Vinik 2003). There is altered expression of neurotrophins and their receptors, as well as neuronal trafficking of neurotrophins in experimental models of DPN. It has been suggested that such changes could in fact contribute to the pathogenesis of DPN, but the data are far from definitive (Fressinaud et al. 2003; Pittenger and Vinik 2003). While large myelinated fibers can be affected in this model, it is the neuropathy associated with the small thinly myelinated fibers and sympathetic neurons that causes the majority of symptoms that adversely affect quality of life (Apfel 1999b). In this context, experimental depletion of NGF or its receptors duplicates many of the neuronal abnormalities observed in this model (Pittenger and Vinik 2003). So perhaps unsurprisingly NGF, which selectively supports the survival of the subset of smaller DRG and sympathetic neurons, was trialled as a potential therapy for DPN due to the consequences associated with degeneration of these neuronal populations (Apfel 2002). A number of animal studies initially demonstrated benefit following NGF treatment for toxic and diabetic peripheral polyneuropathy, which lead to the first clinical trials using neurotrophin-based strategies (Apfel 1999a, b). However, while significant improvement was seen in early trials, studies were discontinued at the phase III stage due to painful side effects (Apfel 2001, 2002). As NT-3 supports the survival of larger DRG neurons, by adopting a similar rationale, it too was trialled as a therapy in animal models of DPN. Utilizing an adenovirus-based vector approach

for the delivery of NT-3 in drug-induced diabetic rats, the slowing of motor nerve conduction velocity and muscle denervation was prevented, suggesting NT-3 retards the development of DPN (Pradat et al. 2001). Combination studies with both NGF and NT-3 in animal models of peripheral neuropathy including DPN suggested that these factors can ameliorate neuronal degeneration (Apfel 1999a). However, as indicated above, this is not a true demyelinating neuropathy, so the lessons learned here in the context of remyelination are unclear.

The contemporary view is that NGF promotes peripheral nerve myelination via activation of neuronal TrkA receptors (Chan et al. 2004). In the adult, no motor neurons and relatively few DRG neurons express TrkA, which tend to be thin and nonmyelinated anyway, so the rationale for utilizing NGF as a strategy to promote peripheral remyelination has limited basis. In fact the pain associated with this clinical trial may be attributable to NGF producing peripheral sensitization of nociceptive DRG neurons through the activation of TrkA. Indeed recent work has demonstrated that increases in peripheral nerve sensitization and pain are due to TrkA activation, p38 phosphorylation, and TRP channel expression (Bron et al. 2003; Obata et al. 2006).

With NGF being an unsuitable therapeutic candidate, what of BDNF? It is certainly implicated in remyelination, as endogenous BDNF has been shown to promote peripheral nerve regeneration and remyelination after sciatic nerve crush injury (Zhang et al. 2000). Electrical stimulation enhanced the remyelination of injured peripheral nerves by inducing BDNF expression and promoting Schwann cell polarization (Wan et al. 2010). However, recent findings have identified that the influence BDNF exerts upon peripheral myelination is more complex, due to BDNF activating distinct receptors: p75NTR to promote myelination and TrkB to inhibit it (Xiao et al. 2009b). On the simple basis of these contrasting influences, BDNF also presents as a less than ideal therapeutic candidate. Nevertheless the identification that BDNF promotes peripheral myelination via neuronal p75NTR is of some interest (Xiao et al. 2009b), as p75NTR has been identified as a global positive regulator of peripheral myelination during development (Cosgaya et al. 2002). In a sciatic nerve injury model in vivo, remyelination was found to be impaired in p75NTR-/- mice (Song et al. 2006). However, it is important to note that approximately 50 % of the DRG neurons are lost in the p75NTR-/- mouse (Murray et al. 1999), complicating this analysis. In a similar sciatic nerve injury model but in nude mice, nerve grafts derived from p75NTR-/- or wild-type mice were transplanted, supplying a source of Schwann cells for remyelination. Analysis of these mice identified that those receiving p75NTR-/- grafts had poorer motor performance, reduced myelin sheath thickness, decreased axonal diameter, and reduced myelin protein expression (Tomita et al. 2007). Collectively these data suggest that p75NTR also plays an important role in the remyelinating process, albeit in a nerve injury model.

Thus, the potential that the neurotrophins hold in promoting peripheral myelination is far from clear, and two issues stand out as requiring attention. The first is the choice of model. Clearly models that utilize nerve injury as the primary pathology are complex, and identifying the selective influence that any putative

therapy exerts on remyelination is confounded. There is a demonstrated need to assess any potential neurotrophin-based therapeutic strategy in a targeted primary demyelinating model, such as experimental autoimmune neuritis (EAN), an autoimmune disease model induced by inoculation with PNS antigen (Gold et al. 2000). The second standout issue is the complexity in the interactions between neurotrophins and their receptors. The cell biology has identified that the neurotrophins exert differing and at times contrasting influences on peripheral myelination, dependent on the complement of neurotrophin receptors expressed by the peripheral neuron. Thus, the therapeutic potential of the neurotrophins per se maybe functionally limited, yet it remains plausible that the selective molecular targeting of p75NTR may, in principle, exert a unified promyelinating response in the PNS. A couple of research groups have recently identified compounds that act as putative selective p75NTR modulators (Massa et al. 2006; Fletcher et al. 2008; Fletcher and Hughes 2009), so the potential of a selective agonist that uniformly promotes the myelination of distinct subsets of peripheral neurons remains a distinct possibility.

#### 3.2 The Neurotrophins Promote Central Remyelination

There are two major groups of CNS demyelinating diseases. The first is the leukodystrophies, a group of genetic disorders that affect the production or metabolism of one of several of the component molecules of myelin. The second is inflammatory demyelinating diseases, caused by autoimmune mediated damage to myelin and oligodendrocytes, with MS being by far the most significant of these diseases. Experimental models of CNS demyelination, including experimental autoimmune encephalomyelitis (EAE) and toxic models of demyelination such as lysolecithin, ethidium bromide, and cuprizone models, have been widely used to examine the therapeutic potential of candidate molecules and to assess the affect they exert upon oligodendroglial and immune cells, as well as their influence on the extent of demyelination and remyelination (Rodriguez 2007).

While the roles that the neurotrophins exert on CNS myelination have been extensively studied in the context of development (Barres et al. 1993; Cellerino et al. 1997; Chan et al. 2004; Xiao et al. 2010), the influence that they exert upon demyelination and remyelination has received substantially less attention. BDNF is the neurotrophin most widely expressed in the CNS and unsurprisingly has been the focus of most attention.

In the normal CNS, BDNF is primarily derived from neurons (Lewin and Barde 1996). However, in inflamed areas in the CNS, such as occurs in MS and EAE, activated astrocytes express increased levels of endogenous BDNF (Stadelmann et al. 2002; Burbach et al. 2004; Linker et al. 2010). Moreover, BDNF is also produced by several immune cell subtypes in vitro, such as activated T cells, B cells, and monocytes (Kerschensteiner et al. 1999; Stadelmann et al. 2002). Thus, BDNF is well placed to exert a modulatory effect upon myelination in this context (Stadelmann et al. 2002). Interestingly in the EAE model, the MS

immune-modulatory drug glatiramer acetate causes a sustained increase in BDNF expression in the brain, which its protective effects are dependent upon (Aharoni et al. 2005; De Santi et al. 2009; Lee et al. 2011).

Several recent studies have implicated BDNF in exerting a functionally protective role following CNS demyelination. In response to the cuprizone model of demyelination, BDNF heterozygote mice exhibit greater demyelination and reduced remyelination in the absence of any change to the total number of oligodendroglia, microglia, astrocytes, or extent of axonal injury (VonDran et al. 2011). In mice in which BDNF has been selectively deleted under the influence of the GFAP promoter, and thus theoretically deleted in oligodendrocytes, astrocytes, and some neurons, a more severe course of EAE and increased axonal loss was observed (Linker et al. 2010). Interestingly, the protective effect of BDNF in this model is temporally restricted, as an inducible conditional knockout strategy demonstrated that BDNF exerts its protective effects primarily during the initial phases of EAE, reducing clinical symptoms and structural damage, whereas deletion of BDNF in later stages of the disease had no significant effect on either disease course or axonal integrity (Lee et al. 2011). This suggests there may be an early window of therapeutic opportunity for the modulation of BDNF levels, at least in EAE.

It appears that not only central nervous system-derived BDNF is important, as mice deficient for BDNF in T cells exhibited progressive disability and enhanced axonal loss in EAE, while mice overexpressing BDNF in T cells exhibited less severe EAE and axonal protection (Linker et al. 2010). Despite this, however, CNS-derived BDNF appears more critical than immune cell-derived BDNF. Experiments utilizing bone marrow chimeras reveal that following CNS deletion of BDNF, immune cell-derived BDNF cannot influence disease severity (Lee et al. 2011). This suggests that while immune cell-derived BDNF can modulate disease, CNS resident cells are the major source of biologically relevant BDNF in autoimmune demyelination. Whether in this context BDNF exerts its influence directly on the myelinated axon/neuron-oligodendrocyte interface remains an open question. In an alternate strategy, treatment with BDNF-engineered bone marrow stem cells also resulted in a significant delay to EAE onset, reduced clinical severity, reduced expression of pro-inflammatory cytokines TNF- $\alpha$  and IFN- $\gamma$ , and increased expression of anti-inflammatory cytokines such as IL-4, IL-10, and IL-11 (Makar et al. 2009). In addition, immune cell expression of TrkB has been shown to play an important role in EAE immunopathogenesis by modulating autoreactive T cell survival and behavior (De Santi et al. 2009). This indicates that BDNF may also influence the immune response, suggesting its protective effect at least in part could be indirect. Ultimately the identity of the cells that BDNF targets for its protective effect may not be absolutely identified until a new strategy is taken, whereupon the EAE model is undertaken in mice in which the BDNF receptor is conditionally targeted in distinct candidate cell types.

In the CNS, TrkB expression is present in various cell types including neurons, oligodendroglial cells, astrocytes, as well as infiltrating immune cells (Cahoy et al. 2008; De Santi et al. 2009), so there is no shortage of potential BDNF targets.

Interestingly, the detection of phosphorylated TrkB was increased in the neurons of EAE-diseased mice compared to the intact axons of control mice, suggesting BDNF is at least exerting a neuronal effect in this context (Linker et al. 2010). As both in vitro and in vivo analyses have identified that oligodendroglial TrkB potentiates CNS myelination (Xiao et al. 2010; Wong et al. 2013), it would be interesting to investigate directly whether oligodendrocyte expression and phosphorylation of TrkB is altered in models of demyelination/remyelination. This would provide some insight into whether the selective promyelinating influence that oligodendroglial TrkB signalling exerts during development is also relevant to remyelination. What is beyond dispute however is that regardless of the model or the strategy to manipulate BDNF or TrkB expression, the effect they exert is a uniformly promyelinating one. Future studies aimed at identifying the cellular and molecular basis of the protective and regenerative effects that BDNF exerts in the context of animal models of demvelinating diseases are critical now in order to establish how these responses might ultimately be targeted and modulated for therapeutic benefit.

# 4 The Therapeutic Potential of Targeting Neurotrophin Receptors for Promoting Remyelination

Despite substantial increases in our understanding of the mechanisms of myelination, demyelination, and remyelination, current therapeutic interventions for demyelinating diseases are suboptimal. These advances in knowledge have yet to be translated into the clinic, meaning that current therapies focus on reducing inflammation rather than directly promoting remyelination. However, as discussed above, there is now a strong body of evidence from preclinical studies pointing to the therapeutic potential of neurotrophins in preventing or slowing demyelination and promoting remyelination.

Human clinical trials assessing the efficacy of human recombinant NGF as a treatment for peripheral neuropathy were undertaken over a decade ago and were discontinued due to side effects including pain (Apfel et al. 1994; Apfel 1999b; Apfel et al. 2000; Apfel 2001, 2002). While the effect on pain sensitization is likely to be an NGF-specific side effect, the complexity of neurotrophin receptor interactions and the more recent identification of the context-dependent influences the neurotrophins exert upon peripheral myelination described above confound the clinical use of recombinant neurotrophins for this purpose. In addition the pharmacokinetic behavior of the neurotrophins is generally not optimal for pharmaceutical use. BDNF, for example, has a half-life in the circulation of less than 1 min in the rat (Poduslo and Curran 1996), and, as one would expect with a protein, there is no strong evidence that it can cross the blood–brain barrier.

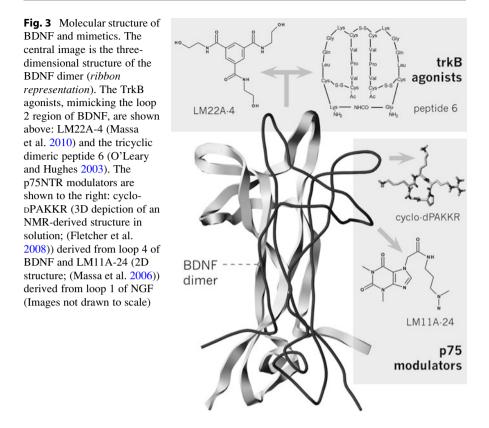
Given the difficulties inherent with using recombinant neurotrophins themselves to treat demyelinating and other CNS disorders, increasing attention has turned to the development of alternative strategies to harness neurotrophic action for clinical use. One means of doing this is to use functional mimetics of the neurotrophins. These compounds would have the advantage of being able to selectively target p75NTR or Trk receptors and could be tailored to have appropriate pharmacokinetic properties. Moreover, exciting recent data describing G protein-coupled receptor (GPCR) transactivation of Trk receptors and their downstream signalling pathways raise the possibility of using an even wider range of small molecules to elicit promyelinating effects.

# 4.1 Putative p75NTR Agonists

Unlike NGF, BDNF does not induce painful side effects and is safe and well tolerated in both animals and humans (Mitsumoto et al. 1994; Ikeda et al. 1995). Based on published data (Chan et al. 2001; Cosgaya et al. 2002; Xiao et al. 2009b), BDNF itself could be considered as a suitable promyelinating candidate to test in the context of peripheral neuropathy. However, as described above, while BDNF promotes myelination of peripheral axons via activation of neuronally expressed p75NTR, it also inhibits the myelination of TrkB-expressing neurons (Xiao et al. 2009b). Thus, at least in the context of peripheral myelination, compounds that selectively activate p75NTR, without concomitantly activating TrkB, would be expected in principle to produce a unified promyelinating response in the periphery.

Recently, two different classes of selective modulators of p75NTR have been described. Structure-based design approaches have been used to obtain conformationally constrained, low molecular weight peptides based on the solvent-exposed loops of BDNF involved in binding to its receptors (Fig. 3). Using this approach, monomeric mimetics of loop 4, exemplified by the cyclic pentapeptide cyclo-[DPro-Ala-Lys-Lys-Arg] (cyclo-DPAKKR), have been developed (Fletcher et al. 2008; Fletcher and Hughes 2009), consisting of the cationic "Lys-Lys-Arg" motif shown to be required for the binding of BDNF to p75NTR (Ryden et al. 1995), presented in a highly conformationally constrained cyclic template. Like BDNF, cyclo-DPAKKR promotes the survival of embryonic chick sensory neurons in culture (Fletcher et al. 2008). However, its mechanism of action is distinct to BDNF, in that it does not cause TrkB autophosphorylation, nor does it lead to phosphorylation of MAPK, a downstream mediator of TrkB signalling, consistent with an effect through p75NTR (Fletcher et al. 2008). In addition, cyclo-DPAKKR has low molecular weight (580) and is highly resistant to proteolysis. These properties of cyclo-DPAKKR and analogues render them unique pharmacological tools to investigate the influence of selective modulation of p75NTR on PNS myelination.

Indeed, recent data indicate the effects that cyclo-DPAKKR exerts upon peripheral myelination are entirely consistent with it acting as a selective p75NTR activator: *promoting* the myelination of distinct subsets of DRG neurons in vitro and requiring neuronal expression of p75NTR. Importantly, local injection of cyclo-DPAKKR adjacent to the developing sciatic nerve in vivo also significantly enhanced myelin protein expression and increased the number of myelinated axons (Xiao et al. 2013). It was also identified that the promyelinating effect of cyclo-DPAKKR was coincident with an increase in the expression of neuregulin-1



type III which provides the key instructive axonal signal that initiates myelination (Michailov et al. 2004; Taveggia et al. 2005). Collectively, these data demonstrate that cyclo-DPAKKR promotes peripheral myelination in vitro and in vivo, suggesting it is a strategy worthy of further investigation for the treatment of peripheral demyelinating diseases.

Recently, Frank Longo's group described a tandem in silico and in vitro approach that leads to the identification of a novel p75NTR ligand, LM11A-24 (Massa et al. 2006). This small molecule, a derivative of caffeine that was selected as a mimic of loop 1 of NGF, has been found to inhibit Aß-induced neurodegeneration and synaptic impairment, block proNGF-induced cell death, and prevent the death of motor neurons in vitro (Pehar et al. 2006; Yang et al. 2008). As a putative selective p75NTR modulator, it would be interesting to investigate the effect that LM11A-24 exerts on Schwann cell myelination in vitro. In particular, it will be intriguing to see if it is able to uniformly promote myelin formation of distinct populations of DRG neurons in vitro and whether this influence can be replicated on peripheral myelin development in vivo. Ultimately, the logical extension of this approach would be to investigate whether both cyclo-DPAKKR and LM11A-24 have the ability to reduce the severity/promote remyelination in an animal model of peripheral nerve demyelination such as EAN.

#### 4.2 Putative TrkB Agonists

In contrast to its apparent inhibitory influence upon PNS myelination, TrkB signalling has been shown to play a key role in promoting CNS myelination. Thus, the selective targeting of TrkB offers a potential means of promoting CNS remyelination. A version of the structure-based design approach has been used to obtain conformationally constrained, low molecular weight peptides that mimic solvent-exposed loop 2 of BDNF, the region involved in binding TrkB (Fig. 3). Using this approach, monomeric mimetics of loop 2 that act as TrkB antagonists (O'Leary and Hughes 1998) and dimeric mimetics of loop 2 that are highly potent putative TrkB agonists (O'Leary and Hughes 2003) have been successfully developed. Like BDNF itself, the putative TrkB agonists promote neuronal survival in cell culture (O'Leary and Hughes 2003). The best of these dimeric peptides, the conformationally constrained tricyclic dimeric peptide 6, is highly potent with an EC50 for the survival of sensory neurons in culture of 11pM (O'Leary and Hughes 2003), that is, the same order of magnitude as BDNF itself, despite being less than one-tenth the molecular weight of the parent protein.

Similarly, an in silico screen with a BDNF pharmacophore followed by lowthroughput in vitro screening in mouse fetal hippocampal neurons led to the discovery of small molecules with nanomolar neurotrophic activity specific to TrkB, typified by LM22A-4 (Massa et al. 2010). LM22A-4 prevents neural cell death with an efficacy equal to BDNF in vitro and causes hippocampal and striatal TrkB activation and improved motor learning in an in vivo model of traumatic brain injury (Massa et al. 2010). It also improves functional recovery and increases neurogenesis following hypoxic-ischemic stroke (Han et al. 2012) and rescues TrkB phosphorylation and improves respiratory function in a model of Rett syndrome (Schmid et al. 2012). Together, these studies demonstrate the potential utility of BDNF – particularly its solvent-exposed loops – as a template for the discovery of novel low molecular compounds that mimic the action of the parent protein.

Adopting a distinct approach, 7,8-dihydroxyflavone was identified in a cellbased screen as a high-affinity TrkB agonist and provokes receptor dimerization, autophosphorylation, and activation of downstream signalling (Jang et al. 2010). Dihydroxyflavone protects neurons from apoptosis in vitro in a TrkB-dependent manner and is neuroprotective in several models of neuronal injury such as kainic acid-induced apoptosis, the transient MCAO occlusion model of stroke, and the MPTP model of Parkinson's disease. It also exerts positive influences in in vivo models of Rett syndrome (Johnson et al. 2012) and depression (Liu et al. 2010, 2012).

The discovery of these small molecules that function as specific TrkB ligands has made modulating selected components of neurotrophin signalling possible for the first time. As the selective modulation of TrkB could achieve more favorable effects than those mediated by BDNF, which also stimulates other neurotrophin receptors, these small molecules provide a potentially powerful therapeutic tool for the treatment of demyelinating disease. The scene is now set to examine the effects that these compounds exert upon oligodendrocyte myelination in vitro and if successful to extend this work to identify whether they exert a protective influence in animal models of CNS demyelination.

# 4.3 Transactivation of Trk Neurotrophin Receptors by GPCR Ligands

One final strategy for the targeting of neurotrophin receptors lies in the discovery that Trk receptors can be transactivated by GPCRs (Lee and Chao 2001). Transactivation of receptor tyrosine kinases by GPCRs is now recognized as an important signalling mechanism allowing the cell to respond to a vast array of extracellular stimuli (Daub et al. 1996; Luttrell et al. 1999; Rajagopal et al. 2004). Both TrkA and TrkB receptors can be activated, in the absence of NGF or BDNF, via a GPCR-dependent mechanism. At least two GPCR ligands, adenosine and pituitary adenylate cyclaseactivating peptide (PACAP), can activate Trk receptors to promote neuronal survival in vitro, an effect dependent upon Trk-mediated activation of Akt activity (Lee and Chao 2001; Lee et al. 2002; Rajagopal et al. 2004). The effect of adenosine and PACAP upon Trk receptor activation is unusual in that it displays very slow kinetics and selectively activates intracellular receptors. Nevertheless, this effect has been shown to be therapeutically relevant in vivo, as transactivation of the TrkB receptor by adenosine agonists promotes motor neuron survival after facial nerve lesioning (Wiese et al. 2007). While studies to date have focussed selectively on neuronal survival, in principle GPCR transactivation of Trk receptors could also exert a promyelinating influence, as not only is the Akt pathway activated by Trk receptors following transactivation but it is also implicated in promoting oligodendrocyte myelination (Lee and Chao 2001; Narayanan et al. 2009; Tyler et al. 2009). Thus, in the context of oligodendrocyte myelination, further studies are clearly warranted to investigate the influence that GPCR transactivation of Trk receptors exerts. However, for Schwann cell myelination the case is less clear, as in vitro studies have revealed that Trk receptor signalling in peripheral neurons exerts contrasting effects upon Schwann cell myelination and the signalling mechanisms that mediate these distinct responses have yet to be elucidated. In totality, a more complete understanding of the potential influences that GPCR transactivation exerts upon models of both central and peripheral myelination in vitro is required in order to establish an evidence-based rationale for future investigations aimed at determining the therapeutic potential of this mechanism of neurotrophin receptor activation.

# 5 Conclusion

It has proven useful to identify the precise influence and the cellular and molecular basis of how the neurotrophins regulate myelination in vitro. Based on these findings, it is now important to attempt to verify these conclusions through the specific conditional targeting of the neurotrophins and their receptors in vivo, in order to develop a clear understanding of the key influences the neurotrophins exert upon myelination. Future studies aimed at identifying whether these mechanisms are relevant not only to development but also in the context animal models of demyelinating diseases will afford specific insight into potential therapeutic approaches that may effect remyelination. The fact that the neurotrophins can exert positive as well as negative influences on myelination in vitro highlights another challenge. However, the evolving development of small molecule neurotrophin mimetics that selectively target distinct neurotrophin receptors affords a degree of hope that ultimately evidence-based rational approaches that enhance these responses will provide therapeutic benefit in the context of human demyelinating disease.

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# Neurotrophic Factors and Ethanol Neurotoxicity

# Margaret I. Davis

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#### Abstract

Neurotrophins are essential for the growth, differentiation, and survival of neurons during development and in the adult. Considerable data have accumulated over the last few decades implicating classical neurotrophins, Nerve Growth Factor (NGF), Brain-Derived Neurotrophic Factor (BDNF), and Neurotrophins 3 and 4 (NT-3/4), in many aspects of ethanol addiction, neurotoxicity, and repair after ethanol withdrawal. Genetic screens in *Drosophila* have identified novel neurotrophic factors and signaling intermediates involved in acute tolerance and pharmacodynamic adaptation to prolonged exposure in rodent. Ethanol modulates neurotrophic factor expression in vivo in a time- and

M.I. Davis

Section on Synaptic Pharmacology, Laboratory for Integrative Neuroscience, National Institute on Alcohol Abuse and Alcoholism, National Institutes of Health, Bethesda, MD, USA e-mail: midavis@mail.nih.gov region-specific fashion. Ethanol inhibits neurotrophin signaling acutely in vivo and in vitro in many brain regions. Conversely, acute and chronic ethanol exposure can upregulate neurotrophin-associated signaling pathways, particularly in brain nuclei associated with anxiety and addiction. Cell death induced by high concentrations of ethanol can be mitigated by exogenous neurotrophins indicating that neurotrophin induction in vivo may also be neuroprotective but ultimately fails over time. Neurotrophin levels in serum and plasma of patients with alcohol use disorders are dynamic and may serve as a surrogate for central nervous system levels. The kinetics suggest that increased levels during withdrawal may be involved in repair, but these analysis are complicated by genetic polymorphisms and the blood component analyzed, particularly with BDNF which is polymorphic in human populations and also produced by platelets. Neurotrophins are intricately involved in pharmacodynamic compensation with prolonged ethanol exposure, addiction-related plasticity, and neurotoxicity, but considerable work remains to be replicated. pharmacological performed and Recent advances targeting neurotrophins and neurotrophin signaling may ultimately be useful for treating ethanol-induced neurodegeneration and aberrant plasticity associated with addiction.

List of Abbr	reviations
AD	Alcohol dependence
ALK	Anaplastic lymphoma kinase
ARND	Alcohol-related neurodevelopmental disorders
BAC	Blood alcohol concentration
BDNF	Brain-derived neurotrophic factor
BFCN	Basal forebrain cholinergic nuclei
cAMP	3'-5' cyclic adenosine monophosphate
ChAT	Choline acyltransferase
E	Embryonic
ERK	Extracellular signal regulated kinase
FAS	Fetal alcohol syndrome
GABA	Gamma amino butyric acid
GDNF	Glial-derived neurotrophic factor
GENSAT	Genetic nervous system Atlas
ICV	Intracerebroventricular
IGF	Insulin-like growth factor
JNK	Jun N-terminal kinase
MAPK	Mitogen-activated protein kinase
MSN	Medium spiny neuron
NAc	Nucleus accumbens
NGF	Nerve growth factor
NMDA	N-Methyl D-aspartate
NT	Neurotrophin
PACAP	Pituitary adenylate cyclase activating peptide

PI3K	Phosphatidyl inositol 3-OH-kinase
PKA	Protein kinase A
PKC	Protein kinase C
PN	Postnatal
RACK	Receptor for activated C kinase
Trk	Tropomyosin-related protein kinase
VTA	Ventral tegmental area

#### 1 Introduction

The majority of heavy drinkers who consume ethanol daily for years to decades show some form of brain pathology either as a result of alcohol-associated neurodegeneration or secondary to nutritional deficiencies, vascular damage, and hepatotoxicity (see Harper 2009; Zahr et al. 2011; Welch 2011; de la Monte et al. 2012 for excellent reviews). Ethanol is even more detrimental to the developing fetus, with Fetal Alcohol Syndrome (FAS) and Alcohol-Related Neurodevelopmental Disorders (ARND) occurring at a combined rate of < 0.7 %in the United States to as high as 5 % in places like South Africa with high per capita alcohol consumption (de Sanctis et al. 2011; May et al. 2009). Chronic and developmental ethanol exposure clearly causes neurodegeneration and cognitive deficits; therefore, it seems intuitive that factors which mediate neuronal survival would be involved in ethanol-induced neurotoxicity. In models of ARND/FAS and in the adult, neurotrophins play a role in cell death, aberrant synaptogenesis, perturbed migration, tolerance, and repair but may also regulate the selective plasticity that occurs with addiction (reviewed by Ghitza et al. 2010; Castren 2004). Whether neurotrophin expression is the result of or induces subsequent changes in synaptic activity is not always clear, but neurotrophins are integral to many ethanol-sensitive processes.

Neurotrophins are a family of proteins initially characterized for the ability to provide trophic support to developing neurons but also regulate synaptic plasticity and survival in mature neurons (reviewed by Reichardt 2006). Nerve Growth Factor (NGF) was the first neurotrophic factor described which supported the growth of chick sensory ganglia neurons (Levi-Montalcini and Cohen 1960). This was followed by isolation and cloning of a family of proteins that includes Brain-Derived Neurotrophic Factor (BDNF), Neurotrophin-3 (NT-3), and NT-4/5 in mammals and their respective Trk receptors (Leibrock et al. 1989; Klein et al. 1989; Lamballe et al. 1991; Maisonpierre et al. 1990b). Since these initial discoveries, many other factors have been isolated that are also trophic for neurons, but these factors do not contain regions of significant homology with the NGF family. Trophic factors explored in alcohol toxicology belonging to the latter group include Insulin-like Growth Factor 1 (IGF-1), Pituitary Adenylate Cyclase Activating Peptide (PACAP), and Glial-Derived Neurotrophic Factor (GDNF).

Classical neurotrophins of the NGF family are cysteine knot proteins that function as dimers. Neurotrophins are expressed as pro-proteins and cleaved to form active proteins, primarily through activity-dependent secretion and protease activation. Neurotrophins can be target-derived, transported trans-synaptically in a retrograde or anterograde direction, or signal through local mechanisms, depending on the brain region and neurotrophin (see Ginty and Segal 2002; Chowdary et al. 2012: Ascano et al. 2012 for reviews). Classical neurotrophins bind to Tropomyosin Related Kinase (Trk) receptors with picomolar affinity. TrkA is the full-length receptor for NGF. TrkB for BDNF and NT-4, and TrkC is the NT-3 receptor, although neurotrophins can bind other Trks with reduced affinity or differential coupling to signaling pathways (Schecterson and Bothwell 2010; Reichardt 2006). Trks also exist as splice variants that result in truncated receptors without signaling capacity, or with only the Shc binding domain (reviewed by Schecterson and Bothwell 2010). Trk receptors can also be transactivated by G protein-coupled receptors, adding further complexity to the signaling repertoire, potential for cell specific signals, and cross-regulation. Activation leads to dimerization and autophosphorylation of tyrosine residues in the full-length Trk receptors, creating docking sites for signaling intermediates (reviewed by Reichardt 2006). Modulation of signaling by ethanol exposure has been demonstrated for many of the major signaling pathways downstream of these trophic factors either individually, sequentially, or in parallel, including Extracellular signal Regulated Kinase (ERK), phosphatidylinositol 3-kinase (PI3K), phospholipase C, Protein Kinase C (PKC), small molecular weight GTPases, calcium, Nuclear Factor (NF) kB, and cyclic AMP (reviewed my Ron and Messing 2013). Neurotrophins are subject to activity-dependent, promoter-specific regulation and therefore regulated by synaptic activity, which is also directly regulated by ethanol exposure. In addition, novel and growth factor-convergent pathways have been identified in Drosophila that regulate ethanol sensitivity, tolerance, and toxicity that remain to be fully explored in mammalian systems (Devineni et al. 2011; Bhandari et al. 2009). Collectively, these data implicate neurotrophins as integral mediators that are regulated by, or regulate, nearly all neuronal processes, albeit with relative selectivity.

Classical neurotrophins also bind p75 with lower affinity, but pro-neurotrophins bind p75 with high affinity (reviewed by Koshimizu et al. 2009). p75 dimerized with sortilin family members to regulate Rho, NF-kB, and Jun N-terminal Kinase (JNK)-p53-Bax pathways, resulting in both survival and apoptosis (reviewed by Friedman 2010). Apoptosis through p75 is mediated by JNK while survival is signaled via NFkB in some cell types (Yeiser et al. 2004). p75 expression declines with age, indicating an active role in development. p75 can also dimerize with Trks to regulate activity, acting as dominant-negative receptors or enhancers of Trk signaling depending on the context, cell type, and neurotrophin concentration (reviewed by Schecterson and Bothwell 2010). For example, intracerebroven-tricular (ICV) injection of pro-NGF induces degeneration of the cholinergic system (Fortress et al. 2011) and mediates hippocampal growth cone collapse through p75-sortilin signaling to Trio-GEF-mediated Rac inhibition (Deinhardt et al. 2011).

Pro-BDNF and cleavage-resistant BDNF induce apoptosis and inhibit migration in cerebellar granule cells (Koshimizu et al. 2009, 2010). Cleavage-resistant pro-BDNF reduces cholinergic hippocampal innervation and hippocampal dendritic spines but does not appear to induce apoptosis in hippocampal neurons (Koshimizu et al. 2009, 2010). Pro-BDNF also mediates long-term depression (Woo et al. 2005). Thus, the maturation state of the neurotrophin, which is activity-regulated, may dramatically influence the receptor selectivity and the subsequent response of the cell.

This chapter will review the background on neurotrophin signaling, expression in animal models of ethanol exposure, and in patients with alcohol dependence. Each section will provide the necessary background information to evaluate the data. While not exhaustive in detail, these sections will provide references to the primary literature and more extensive reviews. Many studies differ in the presentation of dose or concentration and that should be considered. The LD50 for a single intraperitoneal dose of ethanol is  $\sim$ 3.5 g/kg for most mouse strains and range from less than 0.4 g/dL for some rat studies (Prado Carvalho and Izquierdo 1977) to as high as 7–9 g/kg in others (Material Safety Data Sheet). Concentrations are also given in percent (g/dL or mg/dL) or as millimoles per liter depending on the laboratory. Humans become unconscious at Blood Alcohol Concentrations (BACs)  $\sim 0.35$  g/dL, and death is likely at concentrations  $\sim 100$  mM or 0.5 g/dL. Therefore, LD50 should be considered in experimental situations where tolerance is not a factor. Besides the obvious complexities of dose, duration, tolerance, and withdrawal, several experimental caveats in inconsistencies must also be considered when reviewing 30 years of data speckled on a background of exponential scientific discovery and technological advances. These concerns will be addressed as they arise. Current data collectively suggest that work remains to be performed and replicated to gain a better understanding of how temporal changes in neurotrophin expression regulate, and are regulated by, physiological responses within the nervous system and how these influence neurotoxicity, plasticity, and repair after ethanol exposure.

# 2 Neurotrophin and their Receptors after Ethanol Exposure In Vivo: Animal Models

#### 2.1 Nerve Growth Factor

#### 2.1.1 NGF-TrkA Expression and Distribution

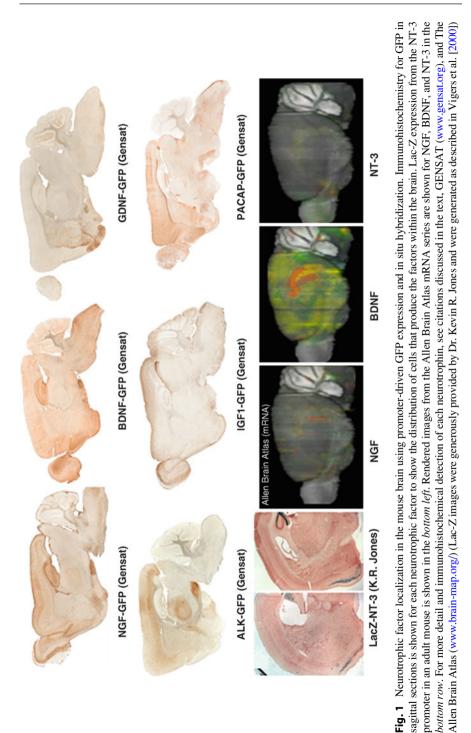
NGF was the first neurotrophin to be examined in models of alcohol-induced neurodegeneration and fetal alcohol syndrome. NGF is expressed in targets of sympathetic neurons and throughout the periphery, with high levels submaxillary gland, heart, prostate, stomach, endocrine system, skin, mast cells, intestine, spleen, lung, skin, and kidney (Korsching and Thoenen 1983; Heumann et al. 1984; Shelton and Reichchardt 1986). Within the CNS, NGF is trophic for Basal Forebrain Cholinergic Neurons (BFCN) where it upregulates choline acyltransferase (ChAT)

activity with an EC50 of  $\sim$ 3 ng/mL (Honegger and Lenoir 1982; Hefti et al. 1984; Mobley et al. 1986; Korsching et al. 1985). Functional studies and developmental expression indicate an essential role for NGF in hippocampal-dependent learning and expression of cholinergic markers in afferent structures (Conner et al. 2009; Conner and Varon 1997; Houeland et al. 2010). NGF also enhances communication between the nucleus basalis and the amygdala (Moises et al. 1995). Functional studies and the expression pattern implicate NGF as a candidate for modulation of ethanol-induced toxicity in the BFCN, hippocampus, and striatum, leading to cognitive deficits and addiction.

NGF mRNA expression (Fig. 1, Allen Institute) and GFP expression from the NGF promoter (Fig. 1, GENSAT) show the general distribution of NGF-producing cells in the adult mouse brain. Expression is sparse in comparison to BDNF shown in the adjacent panels. The highest level of NGF mRNA and protein is detected in the hippocampus within the mossy fiber terminal fields (hilus and stratum lucidum of CA3 and CA2) followed by cortex and olfactory bulbs (Fig. 1, Allen Institute and Gibbs et al. 1989; Maisonpierre et al. 1990a; Conner and Varon 1992; Altar et al. 1997: Conner and Varon 1997). Within the hippocampus, mRNA is localized primarily to interneurons of the hilus, but is also observed scattered in dentate granule cells, scattered interneurons within the CA fields, and interneurons in distal pyramidal cell synaptic fields after colchicine treatment (Ceccatelli et al. 1991). NGF mRNA is also detected in anterior olfactory nucleus, magnocellular preoptic nucleus, nucleus of the horizontal limb of the diagonal band, supramammillary nucleus, striatal interneurons, bed nucleus of the stria terminalis, ventral pallidum, mediodorsal thalamic nucleus, paraventricular hypothalamic nucleus, supraoptic nucleus, lateral and medial septum, substantia innominata, and nucleus basalis. Within the brainstem, NGF is localized to oculomotor, nucleus, facial nucleus, ambiguus nucleus, gigantocellular reticular nucleus, gigantocellular, reticular nucleus (alpha part), inferior olive, reticular nucleus, nucleus of the solitary tract, and vestibular nucleus (Maisonpierre et al. 1990a; Conner and Varon 1992; Altar et al. 1997).

The exact concentration of NGF (and all neurotrophins) in brain regions when measured by ELISA is variable due to extraction procedures, buffer composition, pH, fraction assayed, and the presence or absence of tissue homogenates in the standard curve (see Larkfors et al. 1987; Hoener et al. 1996; Fawcett et al. 1999; Zhang et al. 2000; Soderstom and Ebendal 1995; Whittemore et al. 1986). Calculated value from the studies cited above estimate NGF protein levels at (ng/g wet weight): hippocampus 1.34–6, cortex 0.57–3.5, olfactory bulb 0.85–5.3, basal forebrain 0.672–4.77, cerebellum 0.42–1.86 and striatum 0.16–1.0. Protein levels increase during development in the CNS with a peak that is approximately twice the adult level at postnatal day (PN)14 in cortex, PN14-21 in the striatum and BFCN, and PN7 in the cerebellum (Whittemore et al. 1986; Das et al. 2001; Mobley et al. 1986).

NGF is notoriously sensitive to fixation, pH, and post-fixation times; therefore, many studies purporting to measure NGF by immunohistochemistry may not be measuring NGF. Earlier studies discussed above are replicated in the mouse



expressing GFP from the NGF promoter produced by GENSAT (Fig. 1). NGF immunoreactivity is particularly strong in hippocampal mossy fibers in mature and developing animals with transient production in other dentate lamina during development (Conner et al. 1992; Conner and Varon 1992, 1997). Cells in the stratum radiatum and stratum oriens contain NGF, while a few larger interneurons in the pyramidal cell layer contain NGF immunoreactivity (Conner and Varon 1992; Rocamora et al. 1996; Pascual et al. 1998). Cortical neurons immunoreactive for NGF were generally small (<12 um), but occasional NGF+ cells could be observed that were 13–17  $\mu$ m in diameter, consistent with NGF expression primarily in interneurons (Conner and Varon 1992; Conner et al. 1992; Rocamora et al. 1996).

Regions of neurotrophin expression detected by mRNA frequently do not overlap with the protein distribution measured by immunohistochemistry. This is because NGF is produced in the target region or by local interneurons and retrogradely transported to the cell bodies of the "consumer" neurons (Seiler and Schwab 1984; Conner et al. 1992; Conner and Varon 1992). Colchicine treatment increases somatic immunoreactivity, allowing refined localization of NGF by allowing accumulation of protein in the cells that produce it and blocking retrograde transport to cells that stain for NGF because of uptake from postsynaptic sources. Under these conditions, NGF-expressing cells were detected in the olfactory bulb, tenia tecta, striatum, cortex, septum, BFCN, bed nucleus of the stria terminalis, diagonal band magnocellular preoptic region, ventral pallidum, and substantia innominata. Cholinergic cells with previous immunoreactivity in the septum and BFCN disappeared, thereby segregating the NGF "consumers" from "producers" (Conner and Varon 1992). Thus, many forebrain neurons will contain NGF protein but do not express NGF mRNA, instead obtaining NGF from cortical, hippocampal, and local interneurons (Conner et al. 1992; Conner and Varon 1992; Lauterborn et al. 1995; Whittemore et al. 1986). Similarly, much of the septal NGF is the result of retrograde transport from the hippocampus where it is expressed by cells in the target region for septohippocampal afferents (Conner and Varon 1992; Seiler and Schwab 1984). GFP expression from the NGF promoter also found NGF expression by Fananas astrocytes in the cerebellum and by cerebellar interneurons (Kawaja et al. 2011; GENSAT Fig. 1).

TrkA is widely expressed within the nervous system (see Sobreviela et al. 1994; Altar et al. 1991; Cohen-Cory et al. 1989 for detailed anatomy), including neurons of the striatum, ventral premammillary nucleus, mesencephalic trigeminal nucleus, vestibulocochlear ganglion, prepositus hypoglossal nucleus, raphe nucleus, nucleus ambiguus, glomeruli of the olfactory bulb, area postrema, and nucleus tractus solitarius. TrkA is not present in hippocampal pyramidal cells but is expressed by hippocampal interneurons. The suprachiasmic nucleus, olivary pretectal nucleus, and spinal trigeminal nuclei only contain TrkA immunoreactive fibers. TrkA immunoreactivity is also detected in the leptomeninges, ventricles, and ependymal cells (Gibbs et al. 1989). Iodinated NGF binding is highest in the interpeduncular nucleus and spinal trigeminal tract followed by striatum (higher in lateral striatum), amygdala, diagonal band, nucleus accumbens (NAc), olfactory tubercle, cerebellum, hippocampus, cortex, and subiculum. p75 is notably absent from striatal cholinergic neurons that express TrkA (Sobreviela et al. 1994). Conversely, TrkA is not generally detected in mature cerebellar Purkinje cells that express p75 (Sobreviela et al. 1994; Larkfors et al. 1987, 1996). However, GFP expressed from the TrkA promoter does label a subset of Purkinje cells developmentally, suggesting that some transiently express TrkA. Interestingly, p75 regulates autophagy in Purkinje cells (Florez-McClure et al. 2004). NGF is also trophic for cholinergic interneurons of the striatal matrix in vivo (Van Vulpen and Van Der Kooy 1999) and in vitro (Studer et al. 1994) but has no effect on the survival of striatal medium spiny neuron (Ventimiglia et al. 1995), Purkinje cells (Larkfors et al. 1996), or hippocampal pyramidal cells (Ip et al. 1993).

Genetic deletion of NGF reduces acetylcholine esterase, causes a loss of sympathetic preganglionic neurons, reduces cholinergic septal and striatal neurons, leads to atrophy of BFCNs, and reduces hippocampal innervation by cholinergic afferents (Smeyne et al. 1994; Sanchez-Ortiz et al. 2012), while overexpression of NGF increases the size of septal neurons (Kawaja et al. 1998) and increases cholinergic neuron number and size without altering the innervation of the hippocampus (Fagan et al. 1997; Kawaja et al. 1998). Trk A knockout mice die by 1 month of age, exhibiting severe sensory neuropathies with smaller trigeminal, sympathetic, and dorsal root ganglia (Smeyne et al. 1994).

#### 2.1.2 Ethanol and NGF Expression in Animal Models

NGF levels are sensitive to dose, duration, and withdrawal state, suggesting a dynamic role for NGF (and all neurotrophic factors) in intoxication, tolerance, and toxicity. Ethanol administered acutely via liquid diet causes an increase in NGF and TrkA protein and mRNA in the cerebellum (Wang et al. 2010). Chronic exposure via drinking water also produced a transient increase in NGF in the hippocampus, septum, striatum, cortex, BFCN, and olfactory bulbs (Gericke et al. 2006). After 9 months of exposure, there was a slight reduction in cortical NGF, but no differences in the other regions examined. Shorter exposures (2-4 weeks) of ethanol increased NGF in hippocampus and cortex with a return to control values after 3 months of ethanol exposure (Nakano et al. 1996). Chronic intermittent exposure produced an increase in hippocampal NGF protein levels (Miller 2004), but, in contrast to Gericke and colleagues, chronic exposure for 8 or 24 weeks reduced NGF in most cortical regions cortex and basal forebrain at most time points (Miller and Mooney 2004). Chronic ethanol exposure via liquid diet also reduced TrkA mRNA in the basal forebrain (Miller et al. 2002) and NGF can induce TrkA expression (Holtzman et al. 1992), suggesting failure of NGF to signal with chronic exposure. Combined with reduced basal forebrain TrkA (Miller et al. 2002), this increase in cortical and hippocampal NGF may reflect failure of NGF to transport or a neuroprotective induction that eventually fails with prolonged exposure. In contrast to chronic forced exposure, no differences in striatal NGF levels were observed by McGough and colleagues (2004) after 4 weeks of drinking using the 2 bottle choice model, suggesting some selectivity, depending on the model. The reason for these differences is unclear but may be the result of differential neurotoxicity, the choice to drink, or other environmental factors.

Bruns and Miller (2007) also observed an increase in NGF in the somatosensory cortex in layer V pyramidal cells, not interneurons, with repeated episodic exposure that correlated with the duration of exposure. NGF is generally not generally observed in pyramidal cells (Conner et al. 1992; Conner and Varon 1992; in contrast, see Pitts and Miller 1995; Pitts and Miller 2000), and only pyramidal cells were shown in the micrographs presented by Bruns and Miller; therefore, it is uncertain whether it is ethanol-specific ectopic expression, different pools of NGF, cellular uptake, or resulting from fixation procedures that were significantly different from Conner and colleagues (See Conner 2001 for detailed methods).

While the induction of NGF (i.e., Miller 2004; Gericke et al. 2006; Wang et al. 2010) may correlate with protective capacity and plasticity at early time points, many reports agree that long-term exposure to ethanol decreases NGF levels in the mature rodent hippocampus and cortex (De Simone and Aloe 1993; Aloe et al. 1993; Miller and Mooney 2004; Gericke et al. 2006). Furthermore, reduced hippocampal NGF after prolonged ethanol exposure is associated with atrophy choliner-gic cells in the medial septum; similarly reduced cortical levels are associated with atrophy of septal neurons (De Simone and Aloe 1993; Aloe et al. 1988; Cadete-Leite et al. 2003; Ehrlich et al. 2012). This atrophy and reduced NGF are accompanied by reduced ChAT activity in the hippocampus, septum, and striatum, which can be reversed by administered of NGF via osmotic pump (Cadete-Leite et al. 2003).

NGF-sensitive cells in the developing BFCNs and striatum show transient sensitivity to ethanol that parallel observations in the adult with variable resiliency. Initial studies used chick embryos to show that ethanol was toxic to cholinergic neurons, which was reversed by NGF, and established a window of vulnerability for these neurons (Rahman et al. 1993; Brodie and Vernadakis 1992). A similar window may exist in rodents; a single binge exposure on E15 resulted in a transient increase in hippocampal NGF at PN15 followed by a decline below control values in adulthood and parallel changes in ChAT (Angelucci et al. 1997, 1999).

Rats exposed prenatally to ethanol throughout gestation also have a transient reduction in ChAT activity in the BFCN, striatum, cortex, hippocampus, and septum neonatally (Swanson et al. 1995, 1996; De Simone and Aloe 1993; Fiore et al. 2009a, b). Reversal was associated with an increase in hippocampal NGF (transiently at PN7 and robustly at PN21) and repair in some studies (Heaton et al. 1996). In contrast, others (De Simone and Aloe 1993; Tsuji et al. 2008; Fattori et al. 2008) did not detect any differences in NGF in cortex with prenatal or postnatal exposure but reported a transient decrease in NGF in the hippocampus at PN7 that correlated with reduced cholinergic activity. Similarly, constant gestational exposure to ethanol (but not wine) reduced hippocampal and cortical NGF (Fiore et al. 2009a, b). This was correlated with a reduction in cholinergic neurons in the septum. The survival of cholinergic neurons and NGF induction differs with exposure paradigms, but each outcome is associated changes in NGF levels correlated with cholinergic neuron activity. Whether repair succeeds or fails may depend on the dose and duration of exposure, with binge exposure at a defined window being more toxic to developing BFCNs. As with the pattern in the adult where the ability to upregulate NGF with intermittent exposure may be protective, variability in these data may be the result of the populations that survive the insult and their ability to produce NGF.

NGF levels are acutely increased in the cerebellum in mature mice by ethanol, suggesting a role in adaptation or repair (Wang et al. 2010), but NGF-ethanol interactions have primarily been addressed in the developing cerebellum. Cerebellar Purkinje cells are particularly vulnerable to ethanol toxicity during the first postnatal week but resistant to ethanol-induced cell death during the second postnatal week (West 1993), a period that coincides with the rise to the peak of developmental NGF expression (Das et al. 2001; Maisonpierre et al. 1990a). High dose ethanol (6.6 g/kg/ day, peak BAC 336 mg/dL, cyclic exposure) PN4-10 via artificial rearing did not change NGF levels as detected by ELISA in neonatal cerebellar homogenates but reduced both p75 and TrkA in Purkinje cell dendrites and in cerebellar homogenates (Dohrman et al. 1997; also see Moore et al. 2004a). When this window was further subdivided, Heaton and colleagues (2003) found a transient reduction in NGF only when animals were exposed to ethanol at PN4 (correlating with the developmental period when Purkinje cells are sensitive to ethanol) but an increase at PN7 immediately after exposure and a delayed decrease at 12 h. Surprisingly, exposure on PN7-8 did not change NGF levels, suggesting some form of compensation with prolonged exposure, as was noted in the cholinergic populations. (The levels reported in this study (2003c) are two orders of magnitude greater than previously reported by this lab (i.e., Heaton et al. 2000c) and higher than classical reports. It is possible that this is calculation error, but no correction has been issued.) Furthermore, NGF overexpression from the GFAP promoter rescued Purkinje cells from ethanol toxicity in the neonatal cerebellum without hypertrophy (Heaton et al. 2000b). Purkinje cell number was slightly reduced in the control transgenic mice, suggesting that NGF overexpression alone in a non-activity dependent way may reduce Purkinje cell viability since hypertrophy resulting in fewer neurons per area is excluded but this was not evaluated. NGF overexpression induces aberrant cholinergic sprouting in the cerebellum that might also contribute to the increased survival in this paradigm through cholinergic systems positively coupled to survival pathways and independent of direct NGF effects on Purkinje cells (Kawaja and Crutcher 1997). Furthermore, the locus of NGF action in these studies has not been adequately established and again may suggest robust, dynamic changes in interneurons (Kawaja et al. 2011) or cerebellar afferent fibers since which then modulate Purkinje cell activity. Purkinje cells have not traditionally shown TrkA-mediated survival (Sobreviela et al. 1994; Larkfors et al. 1996) but this does not exclude a role for p75.

Upregulation of NGF at earlier time points after withdrawal or from low levels of ethanol may represent an attempt to compensate for the effect of ethanol on synaptic transmission that fails as producer neurons degenerate. Alternatively, the NGF measured may represent increased pro-NGF, leading to neurodegeneration. In addition, the effect of ethanol on the NGF system specifically in hippocampal and cortical interneurons has not been selectively evaluated in this context, but collectively these data suggest that interneurons may be a central mediator of ethanol effects on NGF-mediated processes, perhaps secondary to reduction in synaptic activity, or death, with prolonged ethanol exposure. As discussed above, neurotrophic activity cannot be examined independent of the chronic and pharmacodynamic effects of ethanol on neural activity and circuit function.

#### 2.1.3 NGF and In Vitro Models of Ethanol Exposure

Cell culture models have provided insight into ethanol pharmacology and toxicology, but many of the in vitro studies discussed below require supraphysiological concentrations of ethanol to induce cell death. These concentrations are generally more than 100 mM or 0.4 g/dL. Many dose–response curves in the studies discussed below range from 0.4 to 6 g/dL and require high concentrations of ethanol to decrease neurite outgrowth and induce cell death. Ethanol impairs neuronal function in vivo at much lower concentrations; therefore, the events leading to cell death in vivo are possibly more subtle, linked to developmental events such as migration and synaptogenesis, may be distinct from the overt toxicity observed over 24–48 h in vitro, and take longer to develop. Therefore, cell death may not be the result of immediate toxicity but may occur secondary to more subtle impairment in the developmental program that leads to cell death at a later developmental stage as the result of aberrant synapse formation or impaired migration. This caveat applies to similar sections on direct toxicity throughout this chapter.

As the prototypic neurotrophin, NGF was the first to be examined for direct modulation of ethanol toxicity in vitro. Chick sympathetic ganglia developed fewer neurites in response to ethanol in combination with NGF in vitro (Dow and Riopelle 1985) and impaired development of NGF-supported structures in ovo that could be reversed by exogenous NGF (Brodie et al. 1991; Brodie and Vernadakis 1992; Rahman et al. 1993). Dow and Riopelle (1985) used ethanol at concentrations ranging from 0.01 to 0.25 g/dL and evaluated neurite formation as a measure of toxicity but found no evidence for cell death at these concentrations. Follow-up in vitro studies replicated the decrease in neurites but extended the concentrations to include doses up to 6 g/dL (Heaton et al. 1993). Ethanol (0.25 g/dL) combined with NGF caused a slight ( $\sim 20$  %) increase in processes in this model but higher concentrations reduced process formation and concentrations above 5 g/dL were required to kill dorsal root ganglia neurons. Similarly, cell death in cultured hippocampal and septal neurons did not occur at concentrations below 2.4 and 1.8 g/dL, respectively (Heaton et al. 1994). NGF was protective against toxicity, even at these extreme concentrations. When cultures were prepared from embryos exposed to ethanol in vivo, they were resistant to cell death at 1.8 g/dL but not 2.4 g/dL (Heaton et al. 1995), suggesting that exposure to ethanol increased resistance to subsequent exposure. In cortical organotypic cultures, NGF has region-specific effects, with neuroprotection only evident within the lower cortical layer developmentally (Mooney and Miller 2007). NGF-mediated neuroprotection is more dramatic in dispersed cultures of cortical, hippocampal, and cerebellar granule neurons (Heaton et al. 2000a; Mitchell et al. 1998; Seabold et al. 1998; Luo et al. 1997; Bonthius et al. 2003), although pyramidal cells and cerebellar granule cells do not respond to NGF through TrkA (Ip et al. 1993; Collazo et al. 1992;

Segal et al. 1992; Barker-Gibb et al. 2001; Courtney et al. 1997; Nonomura et al. 1996; Larkfors et al. 1996). These effects may instead be mediated by p75 in granule cells (Courtney et al. 1997). Alternatively, this may also be due to the relatively high concentration of NGF used, which can nonspecifically activate other Trks. While many neurotrophins are capable of increasing survival in ethanol-exposed cultures, some may be increasing baseline survival without a specific protective effect from ethanol (Bonthius et al. 2003).

PC-12 cells, an immortalized pheochromocytoma cell line with sympathetic neuron-like properties, express TrkA and show enhanced neurite formation, increased microtubule content, and enhanced differentiation in response to 100 mM ethanol ( $\sim$ 0.48 g/dL) and NGF (Wooten and Ewald 1991; Messing et al. 1991: Roivainen et al. 1993: Roivainen et al. 1995). Enhancement was not seen with forskolin and ethanol, but FGF-induced neurite outgrowth was enhanced (Messing et al. 1991). This was accompanied by an increase in Thy-1 expression and subsequently determined to be mediated by PKC- and ERK-dependent pathways (Roivainen et al. 1993, 1995). PKC epsilon, a calcium-independent PKC isoform, was shown to mediate this enhancement of ERK (Hundle et al. 1997), while PKC delta, not epsilon, is responsible for the increase in polymerized microtubule content (Reiter-Funk and Dohrman 2005). Interestingly, PKC epsilon can induce neurite formation independent of catalytic activity (Zeidman et al. 1999) and inactive kinase induces more neurites than wild type PKC epsilon (Ling et al. 2004). Cell cycle arrest and enhanced differentiation were also observed in PC-12 cells, as determined by an increase in neuron-specific enolase expression, a decrease in <sup>3</sup>H-thymidine incorporation, and increased acetylcholine esterase expression with 87 mM ethanol in combination with NGF (Wooten and Ewald 1991). NGF-differentiated PC-12 cells are less sensitive to ethanol-induced cell death in the absence of serum (Oberdoerster and Rabin 1999) and, when specifically induced to differentiate with NGF, became less sensitive to ethanol inhibition of calcium channels (Mullikin-Kilpatrick and Treistman 1995). Therefore, ethanol causes cell cycle arrest and cell death in mitotic PC-12 cells, but once induced to differentiate by NGF, the signaling to ERK is enhanced by ethanol and neurite formation is enhanced, potentially protecting the cell from toxicity but with a potentially detrimental outcome if this were to occur in differentiating neurons.

Recent work using explant cultures of the nucleus basalis suggests more subtle effects than cell death in cholinergic neurons in response to ethanol (Ehrlich et al. 2012). Ethanol reduced ChAT immunoreactivity in neonatal explants without causing cell death or inflammatory cytokine induction which was reversible upon withdrawal. NGF (10 ng/mL) reversed the 100 mM ethanol-induced reduction in ChAT but did not reverse the reduction observed with 50 mM ethanol. Induction of trophic factors with the lower concentration could occlude an NGF effect since NGF was used near the EC50 in these experiments. Perhaps the ability of the cells to produce their own trophic factors is impaired at 100 mM, unmasking the effect. Furthermore, Erlich and colleagues implicate the nitric oxide synthase-p38 MAPK pathway in the reduction in ChAT with both 50 and 100 mM ethanol. These are

intriguing findings and support a role for NGF in repair after prolonged (7 days) ethanol exposure. The time course of NGF inductions varies significantly between studies and duration of ethanol exposure, but collectively, these data suggest induction of NGF that may mitigate some of the toxic effects of ethanol. NGF can be trophic, but if it is induced without concurrent neuronal activity and not cleaved, then pro-NGF can signal cytoskeletal changes or apoptosis through p75. Replicative experiments could focus on determining levels of pro and mature NGF, take advantage of newer technologies to determine the contribution of ethanol-induced changes in NGF and NGF signaling in survival, adaptive plasticity, addiction, and apoptosis.

#### 2.1.4 NGF in Patients with Alcohol Dependence

Genetic polymorphisms and haplotypes in neurotrophins and their receptors in psychiatric and neurodegenerative disorders is an area of active research. The role of neurotrophin polymorphisms in neuropsychiatric disorders is controversial, and many initial association studies were not replicated. Others have reported no association with the underlying disorder but instead indicate a role for genotype in response to treatment (see Table 1). A few definitive associations of neurotrophin mutations with overt neurological disorders have been made, while most associations are weak or only relevant in subsets of patients. This applies to BDNF and other neurotrophic factors as well. Mutations and polymorphisms in NGF or TrkA have been associated with cancer (Scaruffi et al. 1999; George et al. 1998), hypertension (Kapuscinski et al. 1996), Alzheimer's disease (Di Maria et al. 2012; Nagata et al. 2011), anxiety (Lang et al. 2008), white matter structure (Braskie et al. 2012) and congenital insensitivity to pain (Mardy et al. 2001); however, unlike BDNF, no genetic associations with NGF polymorphisms have been reported with alcohol dependence.

Neurotrophins can be measured in the periphery and may be proportional to central levels as well as a surrogate for sympathetic dysfunction and peripheral neuronal activity. Table 1 details measurements of plasma and serum neurotrophin levels in patients with alcohol dependence and during withdrawal. The majority of the reports detailed in Table 1 indicate that ethanol increases peripheral NGF, regardless of the source (plasma vs. serum), in patients still drinking and during early withdrawal (Aloe et al. 1996; Lee et al. 2009; Heberlein et al. 2008; Jockers-Scherubl et al. 2007). NGF levels were found to be an order of magnitude higher in patients that had previously suffered delirium tremens and were not correlated with Korsakoff's Syndrome (Jockers-Scherubl et al. 2007). NGF levels decline toward normal during prolonged withdrawal but remain elevated for as long as 8 months (Aloe et al. 1996; Heberlein et al. 2008; Jockers-Scherubl et al. 2007). Furthermore, downregulation of NGF toward control levels was associated with methylation of the NGF promoter (Heberlein et al. 2011). In contrast, Yoon and colleagues (2006) reported decreased NGF after only 3 months of withdrawal. Ethanol produced a more dramatic decrease in NGF in patients with a family history of alcohol dependence. The patient demographics may contribute to these disparate

Study	Serum/ Plasma	NGF	BDNF	GDNF	Kit
Lee et al. (2009)	Р	112 pg/mL C	861 pg/mL C		DuoSet ELISA
(2009)		137 pg/mL AD <sup>a</sup>	3502 pg/mL AD <sup>a</sup>		R&D
Huang et al. (2008)	S		15.8 ng/mL C		EMax ELISA
			13.9 ng/mL AD		Promega
			15.4 ng/mL 7d WD		
Joe et al.	Р		822 pg/mL C		DuoSet
(2007)			384 pg/mL AD		ELISA
			+FH 247 pg/mL		R&D
			-FH 583 pg/mL		
Yoon et al.	Р	110 pg/mL C			DuoSet
(2006)		71.9 pg/mL AD,			ELISA
		+FH 64.7			R&D
		-FH 83.3			
		>3 mos WD			
Heberlein	Р	19.87 pg/mL C			
et al. (2008)		61.77 pg/mL AD			
		81.8 pg/mL AD <sup>a</sup>			
		21.68 pg/mL 1d WD			
		42.73 pg/mL 7d WD			
Jockers-	S	42 pg/mL			
Scherubl		401 AD no DT			
et al. (2007)		3292 AD + DT			
		225 1d WD			
		108 8d WD			
		144 3 m WD			
		128 8 m WD			
Costa et al.	S		24.2 pg/mL C		EMax
(2011)			23 pg/mL AD		ELISA
			27 pg/mL AD <sup>a</sup>		Promega
			32 pg/mL Abstinent		
Umene-	S	21 ng/mL C			EMax
Nakano et al.		9 ng/mL D			ELISA
(2009)		9.8 ng/mL D + AD			Promega
Meng et al.	S				Chemi-kine
(2011)					Millipore
					(continued)

Study	Serum/ Plasma	NGF	BDNF	GDNF	Kit
Zanardini	S	NOF		ODNF	
et al. $(2011)$			$\frac{\text{S-4.08 ng/mL C}}{\text{S-4.77 ng/mL AD}}$		Quanti-kin
ct al. (2011)	Р		S-4.77 ng/mL AD		R&D
			P-41.7 ng/mL C		
			P-35.97 ng/mL		
			AD upon admission		
Heberlein	S		535 pg/mL C	196 ng/mL C	DuoSet
et al. (2010)			629 pg/mL AD	82.7 pg/mL AD	ELISA
Assoc with			1d WD	1d WD	R&D
thrombocyte			548 pg/mL AD	93.5 pg/mL AD	
count and			7d WD	7d WD	
GGT levels			614 pg/mL AD	81.4 pg/mL AD	_
			14d WD	14d WD	
Huang et al.	S		14.8 ng/mL C		EMax
(2011)			12.3 No DT		ELISA
			6.2 DT		Promega
			13.4 No DT,		
			7d WD		
			8.9 DT, 7d WD		
Aloe et al.	$P^{b}$	32 C			Sandwich
(1996)		37 pg/mL Acute 1 hr			ELISA
		284 pg/mL AD			
		320 pg/mL Late			
		WD			
		239 pg/mL Early WD			
D'Sa (2012)	Р		S-26.6 ng/mL C		DuoSet
	S		S-35.3 ng/mL AD		ELISA
			$P \sim 1.3$ ng/mL C		R&D
			$P \sim 1.4 \text{ ng/mL}$		
			AD		

#### Table 1 (continued)

AD alcohol dependent, C control, D depressed, WD withdrawal, FH family history, d day, m month, S serum, P plasma

<sup>a</sup>Still drinking

<sup>b</sup>Results indicate plasma but methods state "No heparin, EDTA, or protease inhibitors were used"

observations. Patients were European in the first three studies, while the patients were Asian in the Yoon study and the control population varied by 11 years in age. While peripheral sources, concurrent anticonvulsant therapy, and systemic inflammation contribute to serum and plasma NGF levels, these do not appear to be significant covariates when measuring NGF levels in alcohol-dependent patients, nor does the blood component assayed. The well-established association between

stress and peripheral NGF levels (reviewed by Berry et al. 2012) also provides an intriguing link to alcohol dependence and NGF induction. The relative consistency among these studies implies global effects involving elevated NGF in homeostasis, withdrawal hyperexcitability, and repair after prolonged alcohol exposure that parallel observations in rodent models.

#### 2.2 BDNF

#### 2.2.1 BDNF-TrkB Expression and Distribution

While BDNF was the second neurotrophin to be characterized, BDNF distribution and functions are more widespread than NGF. Figure 1 shows the wide distribution of BDNF promoter–driven GFP expression and mRNA in the mature mouse. BDNF expression is regulated by multiple promoters that are differentially methylated and transcribed developmentally, regionally, by activity, pathologically, and upon exposure to xenobiotics (Pruunsild et al. 2007; Aid et al. 2007; Timmusk et al. 1993). BDNF mRNA targeting and translation are also regulated by 5' and 3' UTR sequences (Tongiorgi et al. 2006). BNDF expression level coincides developmentally with synaptogenesis and begins weakly at E13 in the brain (reviewed by Bartkowska et al. 2010). Levels of BDNF remain low until the second postnatal week which coincides with a peak in apoptosis for many structures (Das et al. 2001; Maisonpierre et al. 1990a; Katoh-Semba et al. 1997). BDNF levels are highest in the hippocampus (5.4 ng/g wet weight) followed by hypothalamus (4.33). All other brain regions contain less than 1.5 ng/g wet weight.

BDNF is expressed at high levels by cells in the cortex and hippocampus but is notably absent in the striatum, NAc, and striatal-derived amygdaloid structures (see Maisonpierre et al. 1990a; Ernfors et al. 1990a, b; Conner et al. 1997; Altar et al. 1997 for detailed expression and anatomy). Within the hippocampus, CA3 pyramidal cells and granule cells show the highest mRNA expression with lower expression in CA1 while hilar polymorph cells produce moderate levels of BDNF. BDNF-immunoreactive fibers populate the hippocampal strata oriens, radiatum, lucidum, and lacunosum moleculare, avoiding the stratum pyramidale. High levels of BDNF mRNA are also detected in the thalamus while both immunoreactive fibers and cell bodies containing mRNA are detected in the cortex, septum, BFCN, and the lateral and basal nuclei of the amygdala (Altar et al. 1997; Conner et al. 1997). As with NGF, BDNF is often synthesized in one area and transported, frequently anterograde (Conner et al. 1997; Altar et al. 1997; Li et al. 2012), but also retrograde as is observed in cerebellar granule cells (Bhattacharyya et al. 1997). Notable in addiction research, the medial habenula, striatum, NAc, central amygdala, and bed nucleus of the stria terminalis do not contain significant amounts of BDNF mRNA detected by in situ hybridization or BDNF-immunoreactive cell bodies but contain BDNF-immunoreactive fibers and cells expressing TrkB, consistent with anterograde transport to these (and other) regions (Altar et al. 1997; Conner et al. 1997; Baquet et al. 2004; Li et al. 2012),

The BDNF receptor TrkB is a family of proteins with thirty-six potential splice variants (Luberg et al. 2010), three of which have been well-characterized. TrkB.FL refers to the full-length receptor with tyrosine kinase activity. TrkB-Shc is truncated but retains Shc binding and coupling to the ERK and PI3K pathways. The truncated TrkB.T1 lacks tyrosine kinase activity and was thought to serve as a "decoy" to sequester and inhibit BDNF signaling, but TrkB.T1 coupling to Rho also regulates cytoskeletal dynamics (reviewed by Fenner 2012). TrkB splice variants are also believed to differentially regulate signaling intermediates, with full-length TrkB coupling to transcription through CREB and truncated variants signaling to the cytoskeleton. Both full-length and TrkB.T1 receptors are expressed throughout the brain, but the full-length receptor is exclusively associated with neurons, while TrkB.T1 is expressed by neurons, glia, choroid plexus, and ependymal cells (Altar et al. 1994). Iodinated BDNF and NT-4/5 binding sites were widely distributed similar to in situ hybridization (Altar et al. 1994). The highest levels of TrkB are in hippocampus, cerebellum, and cortex. Diencephalic, hypothalamic, and midbrain monoaminergic neurons express TrkB, and BDNF increases neurotransmitter synthesis and survival in these cells (Altar et al. 1992, 1994; Madhav et al. 2001; Akbarian et al. 2002; Baquet et al. 2004). TrkB is expressed on the majority of mesencephalic dopaminergic neurons, while many of these neurons also produce BDNF (Li et al. 2012; Numan and Seroogy 1999).

Homozygous BDNF knockout mice are not viable and die as neonates. Heterozygous mice exhibit a motor/balance phenotype by 2 weeks of age due to loss of vestibular ganglia (Klein 1994; Ernfors et al. 1994a; Jones et al. 1994). TrkB knockout mice do not eat and also die at birth (Klein et al. 1992). Reduced BDNF impairs taste (Liebl et al. 1997), resulting in abnormal appetitive behaviors in BDNF haploinsufficient mice (Kernie et al. 2000; see Krimm 2007) and, while controlled for and not observed in all experimental cohorts (Hensler et al. 2003; McGough et al. 2004), could contribute to increased alcohol consumption by BDNF haploinsufficient mice in some substrains. Deletion of BDNF also reduces neuropeptide expression in cortical interneurons but does not decrease parvalbumin- and NPY-expressing interneuron numbers (Jones et al. 1994). Recent work has employed region-specific Cre-recombinase-mediated gene deletion strategies and drug-inducible promoters to circumvent the developmental lethality of the full knockout. Combined, both of these strategies validate that deletion of, or reduction in, BDNF impairs hippocampal function, decreases striatal neuron survival, decreases spine density, and induces hyperactivity and hyperphagia (Lyons et al. 1999; Kernie et al. 2000; Rios et al. 2001; Zorner et al. 2003; Gray et al. 2006; Gorski et al. 2003; Li et al. 2012; Baydyuk et al. 2011). Cortex-specific and nigracortex double deletion of BDNF causes degeneration of striatal medium spiny neurons due to the loss of anterograde trophic support (Baquet et al. 2004; Strand et al. 2007; Li et al. 2012). Selective deletion of TrkB from medium spiny neurons impairs motor function, reduces striatal volume, MSN size, spine density, and impairs MSN dendritic development that is accompanied by a reduction in dopaminergic innervation (Baydyuk et al. 2011; Li et al. 2012). Interestingly, a human mutation in the TrkB gene is associated with developmental delay and severe obesity, a phenotype quite similar to the deficient mice (Yeo et al. 2004).

## 2.2.2 BDNF and Ethanol in Animal Models

A detailed review of ethanol and BDNF interactions in addiction and anxiety was recently published (Davis 2008); therefore, this section will concentrate on toxicity and recent controversial reports. BDNF has been extensively studied in the mature brain after acute and chronic ethanol exposure, but the results have varied significantly, often with disparate results in the same species, with the same exposure paradigm and over the same time course. For example, a few studies report no change in hippocampal BDNF levels (Miller et al. 2002; Okamoto et al. 2006), while most find a decrease in hippocampal BDNF with chronic exposure (MacLennan et al. 1995; Miller and Mooney 2004; Tapia-Arancibia et al. 2001; Hauser et al. 2011; Rueda et al. 2012). Still others report a transient increase in hippocampal BDNF with shorter exposure models (Kulkarny et al. 2011; Kalev-Zylinska and During 2007; McGough et al. 2004) and others see increased BDNF during withdrawal (Tapia-Arancibia et al. 2001) or with chronic intermittent exposure (Miller 2004). Disparate results were observed in BDNF mRNA in the hippocampus with intraperitoneal injection, the simplest of exposure paradigms. McGough and colleagues (2004) found an increase at 30 min while Raivio and colleagues (2012) examined levels at 3 h, implicating time as a factor; however, McGough did not observe a change in BDNF mRNA at this time point but found a reduction at 24 and 48 h. Raivio and colleagues conversely did not observe any differences in hippocampal BDNF at 24 h. Much of the discordant data are clustered around the acute time course as most studies agree that doses and exposure times sufficient to cause degeneration decrease BDNF but periods of withdrawal with intermittent exposure may be associated with attempts at repair (Miller 2004). These disparate results are observed in the same brain region and over the same time course, but collectively suggest that BDNF induction eventually fails in the hippocampus with prolonged exposure while upregulation during withdrawal may contribute to excitotoxicity and withdrawal-associated seizures.

Observations are inconclusive in other regions as well. Both increased and decreased BDNF protein levels have been observed in the cortex after chronic ethanol exposure and vary with region and pattern of exposure (Pandey et al. 1999; Miller and Mooney 2004; Miller 2004; Rueda et al. 2012). BDNF protein and mRNA levels in the cerebellum are acutely reduced by ethanol (Wang et al. 2010). BDNF mRNA levels in amygdala and NAc were also acutely reduced in rats after a single intraperitoneal dose of 2.5 g/kg of ethanol (peak BAC 58 mM); levels remained reduced in the frontal cortex at 24 h but returned to control values in the amygdala and hippocampus. The acute phase of this response is consistent with a decrease in excitatory transmission. BDNF mRNA in the NAc was initially reduced but increased in the ventral tegmental area (VTA) at 3 h, rebounding to increase at 24 h in both structures (Raivio et al. 2012). Kerns and colleagues (2005) also observed an increase in BDNF in NAc by microarray with acute exposure that was mouse strain specific. The acute increase in the VTA could result from direct excitation, reduced inhibition, or accumulation of mRNA that fails to transport but suggest that the VTA may be a source for accumbal BDNF mRNA levels since cortical levels remained lower. Microarray analysis also identified BDNF as a gene with reduced expression in the prefrontal cortex after chronic intermittent ethanol exposure (Melendez et al. 2012). Interestingly, the array analysis also detected increased Oprl1 and extensive modulation of mRNA species belonging to the actin dynamics and MAPK signaling nodes consistent with a pharmacodynamic process. Consistent with a decrease in cortical BDNF mRNA, Bosse and Mathews (2011) measured BDNF protein levels in the striatum after acute ethanol exposure in both wild type and BDNF haploinsufficient mice. Both exhibited a decrease in BDNF; furthermore, the deficient mice did not respond to ethanol with a robust increase in dopamine when measured by in vivo microdialysis.

This is in contrast to work from the Ron/Janak laboratory that did not detect changes in cortical or NAc BDNF with acute exposure or self-administration (Logrip et al. 2009; McGough et al. 2004). Hippocampal BDNF was also elevated by ethanol through a Receptor for Activated C Kinase (RACK1) dependent pathway after both acute and prolonged (4 weeks) ethanol exposure (McGough et al. 2004; Jeanblanc et al. 2009, 2006; He et al. 2010). The data contrast the observations discussed above in extended striatum, implicating an acute increase in dorsolateral striatal BDNF that is negatively correlated with ethanol consumption (McGough et al. 2004; Logrip et al. 2009; Jeanblanc et al. 2009). This is supported by the observation that BDNF haploinsufficient mice voluntarily consume more ethanol than wild type litter mates, but these cohorts do not exhibit generalized hyperphagia, as discussed above (Hensler et al. 2003; McGough et al. 2004). Furthermore, viral mediated expression of BDNF siRNA in the dorsolateral striatum clearly transduced medium spiny neurons identified with GFP and replicated the phenotype of increased drinking (Jeanblanc et al. 2009). RACK1 mediated the ethanol-induced BDNF expression since Tat-RACK1 administration increased BDNF, reduced ethanol intake through TrkB, and reversed the phenotype in haploinsufficient mice when administered in the dorsolateral striatum (McGough et al. 2004; Jeanblanc et al. 2006). BDNF induction by ethanol was also observed in vitro with immunofluorescence in striatal neurons (Logrip et al. 2008), neuroblastoma cells (He et al. 2010), and in hippocampal pyramidal cells (McGough et al. 2004; He et al. 2010), requiring RACK1 for gene expression in each cell type. RACK1 activation of BDNF transcription is purported to be mediated by cAMP (Yaka et al. 2003b, c; He et al. 2010; Neasta et al. 2012), 14-3-3 association (Neasta et al. 2012) and epigenetic modulation of BDNF promoter IV (He et al. 2010). Interestingly, cAMP signaling via PACAP also appears to use this pathway to induce BDNF (Yaka et al. 2003a, b). Cyclic AMP causes dissociation of RACK1 from the NMDA/Fyn complex, thereby increasing Fyn phosphorylation of the NMDA receptor and increased NMDA activity. Collectively, these data place cAMP upstream of many key events involved in adaptation to ethanol exposure but implicate RACK1 and not CREB as the key transacting factor in this pathway.

Similarly, BDNF induction was observed in MSNs of the central and medial amygdala by Pandey and colleagues, where interference with BDNF expression using antisense oligonucleotides increased anxiety-like behavior and ethanol consumption (Pandey et al. 2005). Furthermore, alcohol-preferring rats produced less central amygdala and BNST BDNF mRNA and protein and exhibited higher

anxiety-like behavior (Moonat et al. 2011; Prakash et al. 2008). In this case, CREB appears to be a key intermediate in the behavioral response but Pandey and colleagues place CREB and ERK downstream of BDNF induction. While Pandey's group did not detect strain differences in BDNF in the NAc, Yan and colleagues found a reduction in accumbal BDNF in the preferring strain (Yan et al. 2005). Together, these data implicate reduced BDNF in the basal ganglia in alcohol appetite, but the anatomical locus is not resolved. As with MSNs in the striatum, BDNF protein should be delivered to the central amygdala via afferents (Conner et al. 1997) but the source of the mRNA and efficacy of BDNF siRNA and antisense oligonucleotides were unequivocally shown to be locally mediated within medium spiny neurons in the dorsal striatum and striatal-like nuclei of the amygdala. This was not concluded by Riovio (2012) and colleagues whose data are difficult to reconcile and will require further anatomical, pharmacological, and molecular investigations.

Variable results have also been reported in developing animals with BDNF expression, likely amplified by the low levels of BDNF in neonates and stress. Many fetal alcohol experiments focus on the developing cerebellum because the ontogeny of specific neuronal populations, their migration, synaptogenesis, and the factors modulating these processes have been well-characterized (reviewed by Sotelo 2004). The majority of studies examining BDNF levels, TrkB, or BDNF signaling have shown that ethanol inhibits the neurotrophic activity of BDNF in the cerebellum, and this may contribute to cerebellar apoptosis and impaired foliation. Ge and colleagues (2004) performed a detailed dose-response, time-course analysis and found a rapid decrease in BDNF and TrkB mRNA in the cerebellum after exposure on PN4 but not PN9, suggesting a role for BDNF in the selective vulnerability at this time point when levels are low. In a follow-up study (Light et al. 2002), TrkB was detected in cerebellar Purkinje cells during the vulnerable first postnatal week; ethanol decreased TrkB immunoreactivity as well as TrkB.T1 and TrkB mRNA in these experiments. Moore and colleagues found an increase in TrkB in the cerebellum of females exposed prenatally but no other alterations in cerebellar TrkB with prenatal or postnatal exposure (Moore et al. 2004a, b). Moreover, Heaton and colleagues did not observe significant regulation of BDNF protein levels in cerebellum with numerous exposure windows that span postnatal development but subsequently reported a transient increase in BDNF immediately after exposure on PN4 that normalized by 2 h (Heaton et al. 1999, 2003a). Exposure on PN7 increased BDNF levels at 2 h and decreased levels at 12 h in this study. Observations made in juvenile rats after a single binge of ethanol indicate no changes in cerebellar BDNF mRNA (Kulkarny et al. 2011), suggesting modulation may be more dynamic in younger animals.

Multiple lines of evidence support a role for BDNF expression and signaling in the cortex and hippocampus in the development of FASD, but the sequence and mechanisms have not been fully resolved. A few reports have suggested an increase in BDNF (and related TrkB agonist NT-4) immediately following exposure. Increased BDNF/NT-4 levels were measured in combined "cortex/striatum" at PN1 after prenatal exposure and at PN10 after PN4 -10 (Heaton et al. 2000c). Surprisingly there was no significant developmental increase detected in BDNF between PN10 and PN21. By contrast, Heaton and colleagues (2003a) reported bidirectional changes in BDNF cortex after exposure on PN7 and increased BDNF on PN10 after neonatal exposure that resolved by PN21. It should be noted that the levels of BDNF measured in these assays were very low (50–80 pg/g wet weight) and may have been below linear detection with the dilutions given in the text (100-fold). Others have found that fetal exposure decreases BDNF in the cortex (Climent et al. 2002; Fattori et al. 2008; Feng et al. 2005; Caldwell et al. 2008). In addition, Caldwell and colleagues (2008) found decreased BDNF and specific decreases in transcripts III, IV, and VI in the medial frontal cortex and hippocampus.

Similar differences were observed in the hippocampus where Heaton and colleagues (2000c) find a transient increase in hippocampal BDNF at PN10 with neonatal exposure and no change with prenatal exposure. However, Fattori found a decrease in BDNF after PN5-8 exposure. Feng and colleagues (2005) and Caldwell colleagues (2008) also report a decrease in BDNF after gestational exposure. Miki et al. (2008) exposed animals from PN10-15 via vapor inhalation and also noted a transient increase in hippocampal BDNF mRNA from PN16-PN30 that varied in duration depending upon the control group comparison. These observations were less robust in older animals and complicated by differential BDNF expression between the maternally reared control group and the separation control group such that BDNF levels were reduced when compared to the separation control animals but unchanged relative to the maternally reared animals at 60 days of age (Miki et al. 2008). Similarly, separation of PN5 pups increased PKC activity in the hippocampus, which correlated with increased ERK signaling (Davis et al. 1999). These observations illustrate how stress significantly affects signaling in the developing brain and may be another variable contributing to the differential effects of ethanol on neurotrophin levels and signaling between laboratories and exposure paradigms. While it is difficult to draw firm conclusions from the current data, combined, these studies suggest that ethanol may acutely and transiently increase BDNF in some brain regions developmentally. Correlations with cell death suggest that the increase may reflect an attempt at repair, but may ultimately lead to a reduction in BDNF production in the mature structure if administered during a vulnerable window.

Ethanol also produces variable effects on TrkB that may be dependent not only on timing but also on gender (Moore et al. 2004a, b). However, in spite of this overwhelming confound, others have found that prenatal exposure increases TrkB/ TrkB-T1 ratios as neonates (Climent et al. 2002). Baseline neurotrophin levels and BDNF levels are responsive to multiple environmental and experimental factors that can make accurate measurements difficult. More experiments examining acute effects on signaling during exposure and compensatory regulation over time and after in specific cell types are required. It is also unclear whether these changes result in a decrease in activity-dependent BDNF release or represent differences in synaptic complexity. Examination of BDNF signaling (discussed below) suggests that ethanol causes a decrease in the ability of TrkB to transduce signal and desensitize, which may lead to compensatory regulation of both BDNF and TrkB in vivo. There is considerable data to indicate a role for aberrant BDNF modulation during post-exposure development, but this may be secondary to impaired synaptogenesis, activity, and neuronal apoptosis (Bhave et al. 1999; Feng et al. 2005; Miki et al. 2008; Caldwell et al. 2008; Climent et al. 2002).

#### 2.2.3 BDNF and In Vitro Models of Ethanol Exposure

As with NGF in the previous section, BDNF diminishes ethanol-induced hippocampal and cerebellar granule cell death in vitro (Heaton et al. 2000b; Bonthius et al. 2003). The neuroprotective effects of NMDA in cerebellar granule neurons are also mediated by BDNF induction (Marini et al. 1998; Bhave et al. 1999) and likely involve PI3K since the survival-promoting effects of BDNF were blocked by PI3K inhibition (Bhave et al. 1999). Ethanol also decreased Akt phosphorylation in the developing, but not adult, brain (Chandler and Sutton 2005), implicating PI3K/Akt in the selective vulnerability of the developing brain to ethanol.

Neurotrophins also reverse many of the effects of ethanol on synaptic activity. Exogenous BDNF reverses ethanol attenuation of GABA-A responses in cerebellar granule cells (Ericson et al. 2003). Ethanol also reverses BDNF-mediated long-term plasticity at developing hippocampal CA3 GABAergic synapses through inhibition of voltage-gated calcium channels (Zucca and Valenzuela 2010). At excitatory synapses, ethanol blocks BDNF enhancement of NMDA (NR2B-containing), but not AMPA, responses in hippocampal neurons (Kolb et al. 2005). As a control, the authors show that NGF and NT3 do not alter synaptic transmission, indicating cell type selectivity in the ion channels modulated by specific neurotrophins as would be expected from the anatomical distribution discussed above. Kolb and colleagues further dissected the effect of ethanol on the channel from the BDNF enhancement by using ethanol concentrations that did not have an impact on NMDA signaling, indicating that the effect of ethanol is independent of NMDA receptor inhibition.

In contrast to the experiments discussed above where chronic ethanol enhanced NGF-stimulated ERK activity in PC-12 cells, acute and chronic ethanol administration decreases activation of ERK in primary neurons and in the rodent brain (Davis et al. 1999; Kalluri and Ticku 2002, 2003; Tsuji et al. 2003; Chandler and Sutton 2005; Han et al. 2006; Ohrtman et al. 2006; Fattori et al. 2008 however, see Chen and Charness 2012 and Acquaah-Mensah et al. 2001). Regulation of ERK by ethanol also contributes to reduction in hippocampal LTP (Roberto et al. 2003). Ethanol acutely reduces TrkB signaling to ERK in cerebellar granule cells (Li et al. 2004; Ohrtman et al. 2006), but does not change TrkB phosphorylation state (Li et al. 2004), indicating an intracellular site of action downstream of the receptor. Neonatal hippocampal slices stimulated with BDNF displayed reduced nuclear translocation of phosphorylated ERK in hippocampal pyramidal cells in CA1 (Davis et al. 1999). Inhibition of BDNF-stimulated ERK phosphorylation by ethanol in cerebellar granule cells is dependent upon BDNF concentration and not modified by NMDA receptor antagonists (Ohrtman et al. 2006). Low concentrations of ethanol inhibit ERK activation only when BDNF was present at concentrations in the rising portion of the high affinity concentrationresponse curve (<5 ng/mL), while high concentrations of ethanol also block the desensitization of the ERK response at longer times (30-60 min), and at high BDNF concentrations (>10 ng/mL). Thus, there was no apparent inhibition with 50 ng/mL of BDNF after 20 min, but the peak was never reached in the ethanol-exposed cells while the control curve is declining (Ohrtman et al. 2006). This was also reported recently by Chen and Charness (2012) who did not observe ethanol inhibition of BDNF-stimulated ERK activity with 50 ng/mL of BDNF, but this study did not perform the same pharmacological analysis as Ohrtman and colleagues. Both PKA and PKC regulate ERK, and reduced ERK activity correlates inversely with PKA activity and positively with a concurrent decrease in calcium-sensitive protein kinase C activity in the developing hippocampus (Davis et al. 1999); therefore, acute inhibition of TrkB signaling to ERK may be independent of Trk activation and ultimately cross-regulated by other pathways with different concentrations of ethanol.

In addition to mediating survival, BDNF also regulates migration and synaptogenic processes. Mutations that either upregulate or downregulate signaling through growth factor pathways and their associated GTPases cause a myriad of neurodevelopmental disorders resulting from accelerated or decelerated development (Stornetta and Zhu 2011). Perturbation of signaling that results in subtle changes in the rate of migration and synaptogenesis might also contributes to the development of alcohol-related neurodevelopmental disorders without inducing overt cell death. Numerous reports indicate that concentrations of ethanol as low as 25 mM inhibit neuronal migration neurite outgrowth through the adhesion molecule L1 (Charness et al. 1994; Bearer et al. 1999), and inhibition of ERK and Src and implicated in inhibition of L1-mediated neurite formation by ethanol (Tang et al. 2006; Yearney et al. 2009). In contrast, a few studies report increased migration and increased neurite formation in response to ethanol, as discussed above, and increased basal spines on pyramidal cells treated with chronic intermittent ethanol (Kroener et al. 2012). Developmentally, Powrozek and Olson (2012) observed increased dendrite formation in cortical explant cultures treated with ethanol. Proliferating neuroepithelial precursors exposed to ethanol display a differentiation-associated increase in migration (Camarillo and Miranda 2008) and ethanol increased the rate of axon formation in hippocampal pyramidal cells (Lindsley et al. 2011). This overall increase in axon growth rate was associated with ethanol inhibition of BDNF-stimulated Rac1/Cdc42 activation and increased Rho activity in hippocampal axonal growth cones, leading to increased growth cone surface area (Lindsley et al. 2011). Although ethanol increased the surface area of growth cones, the levels of active Rho GTPases in axonal growth cones were not affected in the absence of exogenous BDNF. These observations are in contrast to a recent study from Chen and Charness (2012), detailing reduced axon growth that was not mediated by BDNF and was dependent on Src, albeit with different substrates. We have also observed an increase in BDNF-stimulated migration/dendrite formation in cerebellar granule cells using a Boyden chamber assay (Hassoun et al. 2007a; See Hassoun et al. 2007b for detailed methods). As discussed above, ethanol also enhances differentiation in some cells, therefore cell cycle arrest, possible through ERK inhibition in cycling cells, and premature differentiation would increase the pool of migration-competent precursors if administered during the correct developmental window. This does not necessarily have to increase the rate of migration, which has been shown to be reduced in the cerebellum after ethanol exposure (Jiang et al. 2008), but may simply increase the number of cells that migrate toward BDNF.

Oxidative stress, also known to modulate neurotrophin signaling, has recently been implicated in the etiology of some forms of autism (Rossignol and Frye 2012; Sajdel-Sulkowska et al. 2011) as well as the development of FAS/ARND (Brocardo et al. 2011). A target that has been suggested to mediate the toxic effects of ethanol is cJun N-terminal Kinase (JNK), a member of the mitogen-activated protein kinase family, which is induced by ethanol, genotoxicity, and oxidative stress (reviewed by Weston and Davis 2007). p75 also activates the JNK pathway (Koshimizu et al. 2010). JNK, as the name implies, phosphorylates c-Jun, a member of the AP-1 (Fos/Jun) transcription factor complex. "JNK" represents 3 families and as many as 16 splice variant isozymes that do not always correlate with molecular weight but is generally divided into 3 molecular weight categories of 58, 54, and 46 kDa. Suppression of Akt activates JNK (Shimoke et al. 1999), but apoptosis in cerebellar granule cells induced by growth factor withdrawal, glutamate, and beta amyloid is not mediated by JNK (Gunn-Moore and Tayare 1998). Some reports suggest that JNK with an apparent molecular weight near 60 kDa is involved in ethanol-induced apoptosis and ethanolinduced oxidative stress, but only one isoform was observed by western blot (Heaton et al. 2003a, 2012). Han and colleagues (2006) observed ethanol-induced increases in p46 and p54 phosphorylation that are associated with apoptosis in PN7 rat pups. However, when phosphorylated cJun is measured as a surrogate for JNK activity after ethanol exposure, there is no overlap between cells undergoing apoptosis and phospho-cJun in immunostained brain sections (Young et al. 2008). Moreover, JNK activation induces genes associated with protection in Drosophila in response to oxidative stress (Wang et al. 2003) and mutations that enhance JNK signaling in Drosophila can extend lifespan (Karpac and Jasper 2009). BDNF stimulation of AP1 DNA binding activity is also reduced by ethanol in cerebellar granule cells (Li et al. 2004). The effects of JNK activation may therefore be isozyme or splice variant, specific as these data suggest reduced genomic signaling through Fos/Jun after ethanol in cerebellar granule cells and a non-genomic, possibly protective, isoform-specific, role for JNK in ethanol toxicity which has not been explored.

The JNK pathway stimulates both dendrite and axon formation through phosphorylation of proteins such as MARKS and doublecortin that are associated with cytoskeletal dynamics (Oliva et al. 2006; Bjorkblom et al. 2012; Jin et al. 2010). JNK1 is associated with microtubules where it regulates axodendritic length (Tararuk et al. 2006), and genetic deletion of JNK or intermediates in the JNK pathway leads to axonal growth defects (Chang et al. 2003; Koushika 2008). Immunohistochemical staining for phospho-JNK in the neonatal mouse brain labels primarily developing axon pathways and not apoptotic cell populations (Davis, unpublished observations). A role for JNK regulating migration and axon outgrowth in response to ethanol is further supported by the observations that ethanol increases integrin and NCAM levels (Vangipuram et al. 2008; Miller et al. 2006), increases laminins (Vangipuram et al. 2008), increases Fyn activity (Yaka et al. 2003b), increases PKC, and increases FAK phosphorylation in cerebellar granule cells (Sivaswamy et al. 2010), which all regulate cytoskeletal dynamics (Becchetti and Arcangeli 2010). JNK may therefore be an overlooked contributor to ethanol effects on the neuronal cytoskeleton and may serve protective as well as apoptotic functions.

# 2.2.4 BDNF in Patients with Alcohol Dependence

Variability in peripheral BDNF levels in alcohol-dependent patients is more extreme than NGF, making it difficult to deduce trends from the current data. The levels of BDNF reported in Table 1 are variable, often with standard deviations greater than the mean. Stress, depression, hematologic parameters, xenobiotics, age, and gender are all known to influence peripheral BDNF levels (reviewed by Autry and Monteggia 2012; Lommatzsch et al. 2005). BDNF levels detailed in Table 1 do not correlate with the source (plasma vs. serum), method used, or duration of abstinence. The source of the variability could be innate, due to simultaneous use of anticonvulsants during withdrawal, or due to sample preparation. Many experiments used serum, which includes BDNF released by thrombocytes, and is orders of magnitude higher than the plasma concentration (Lommatzsch et al. 2005). Two studies state different sources in the methods compared to the results section, although the levels reported suggest the source (ng vs. pg concentrations). In other studies, it is not clear that the centrifugation steps are sufficient to remove the thrombocytes, while others use sedimentation forces likely to rupture the cells. Zanardini and colleagues (2011) and D'Sa and colleagues (2012) directly compared serum and plasma BDNF. The first study reported a small increase in both components, while the second study only detected an increase in BDNF in serum in alcohol-dependent patients. Heberlein and colleagues (2010) also addressed this confound by simultaneously measuring thrombocyte activity and serum BDNF levels, observing a positive correlation between serum BDNF levels and both gamma glutamyl transferase levels and thrombocyte function. In addition, Costa and colleagues (2011) found a nonsignificant but suggestive association of BDNF levels with platelet count. Alcohol dependence and withdrawal are known to alter thrombocyte levels and coagulation cascades with robust rebound upon withdrawal (Schmitt et al. 1999; Chanarin 1982). BDNF is also produced and released by the vascular endothelium (Nakahashi et al. 2000), further complicating any correlation between central function and peripheral levels in patients with underlying cardiovascular disease.

Peripheral levels may provide valuable information since the production and release mechanisms regulating plasma and serum levels are likely to be similar to mechanisms governing central production and release, even if they are not the same pools of BDNF. Interestingly, this was predicted by Chinarin in 1982: "After platelet aggregation the platelets contract and squeeze various metabolites out of the platelet; this is called the 'release reaction'. This phase is also curtailed or even absent in platelets from alcoholics". BDNF polymorphisms also have the potential to regulate peripheral levels, but these reports are similarly variable and primarily examine serum BDNF. The Met allele of the Val66Met BDNF polymorphism decreases neuronal BDNF secretion (Egan et al. 2003; Chen et al. 2004, 2006). Some evidence supports an association between the homozygosity for the Met or

Val allele with lower or higher BDNF levels, respectively (Ozan et al. 2010; Cirulli et al. 2011). In contrast, no association between genotype and serum levels were found in the majority of patient populations examined thus far (Yoshimura et al. 2011; Trajkovska et al. 2007; Terracciano et al. 2011) and, paradoxically, patients with the Met allele were even found to have higher serum BDNF concentrations (Lang et al. 2009). Thus, factors other than genotype may be responsible for the majority of the variability in serum and plasma BDNF.

BDNF is highly polymorphic and linkage analysis identified an alcoholism susceptibility gene that mapped near the BDNF locus, 11p13 (Uhl et al. 2001). The human BDNF gene is encoded by 11 exons and 9 promoters (Pruunsild et al. 2007) and is polymorphic between different human populations, further complicating association analysis (Petryshen et al. 2010). Three polymorphisms have been examined in alcohol-dependent patients; these polymorphisms change BDNF secretion and activity-dependent processing (Egan et al. 2003; Chen et al. 2004, 2006). The first is a missense change (G196A, rs6265) that results in a valine to methionine substitution (Val66Met). The Met allele shows impaired trafficking and activity-dependent secretion, and humans homozygous for the Met allele have reduced hippocampal and amygdalar volume, increased anxiety, increased heart rate, and poor working memory (Egan et al. 2003; Hariri et al. 2003). Humans homozygous for the Met allele have recently been reported to have higher anxiety, increased cortisol, and consume more alcohol (Colzato et al. 2011). Abstinent Met/Val and Val/Val patients differ in their rate and locus of recovery measured by MRI, with Val/Val homozygotes showing more gray matter recovery while heterozygotes had more recovery in white matter measures (Mon et al. 2012). Transgenic mice carrying the human Met allele and their human counterparts were also impaired in models of extinction learning, with human Met/Met homozygotes demonstrating impaired fronto-amygdaloid activity by functional magnetic resonance imaging (Soliman et al. 2010). A second polymorphism (C270T) is present in the 5' noncoding region that can regulate transcription or translation. The third genotype examined is a G712A polymorphism that has not been extensively investigated molecularly but may be associated with Parkinson's Disease (Chen et al. 2011). Initial studies suggested BDNF polymorphisms were associated with psychiatric disorders; however, subsequent studies and meta-analysis have failed to support associations for most major neuropsychiatric disorders, while studies contain too few patients to asses validity in less common disorders at this point (reviewed by Hong et al. 2011).

Anxiety is a precipitating factor for alcohol abuse. Association between the Met allele and higher anxiety-like behavior has been observed in both humans and transgenic mice expressing the human Met allele (Jiang et al. 2005; Montag et al. 2010; Chen et al. 2006). BDNF polymorphisms have been examined by several groups and the associations with alcohol dependence have been weak and variable in spite of promising initial observations. Matsushita et al. 2004 found that G196A (Val66Met) BDNF polymorphism frequency did not differ significantly from controls but alcohol-dependent patients with violent tendencies and a history of delirium tremens had an increased frequency of Met/Met genotype. Antisocial

alcohol behavior and alcohol dependence may also segregate with a TrkB variant (Xu et al. 2007). The Met allele was also associated with earlier disease onset in this study. Met carriers were subsequently shown to consume more alcohol than Val66Val or Val66Met patients which segregated with alcohol use disorders (Shin et al. 2010; Colzato et al. 2011). Reduction of BDNF is associated with aggression in mice (Chen et al. 2006), but the genetic association between aggression, Met/Met, and alcoholism was not observed in the majority of replicate analyses (Tsai et al. 2005; Muschler et al. 2011; Wojnar et al. 2009; Grzywacz et al. 2010). In a subgroup analysis, the Met allele was overrepresented in alcoholdependent patients with depression compared to controls or dependent patients without depression (Su et al. 2011). While no association was found with dependence, Woinar and colleagues (2009) found an association between the Val66Val genotype and relapse that was greater when restricted to analysis of those with a positive family history of alcohol use disorders. In support of this phenotypic association, the Val/Val is also associated with impulsivity (Gatt et al. 2009), which can contribute to relapse. A modest association between SNP G712A and substance dependence was also described but did not specifically examine the alcoholdependent subgroup (Zhang et al. 2006). These data do not suggest a strong association between the BDNF polymprphisms and alcohol dependence but may identify a subgroup of patients and indicate response to treatment. Interestingly, over 60 polymorphisms in the BDNF gene have been identified to date, so lack of association at two loci does not exclude a role for BDNF genotype as a modifying factor in alcohol abuse, dependence, and toxicity.

# 2.3 NT-3

#### 2.3.1 NT-3-TrkC Expression and Distribution

NT-3 is widely expressed in the periphery, with high levels in the kidney, spleen, heart, and muscle (Maisonpierre et al. 1990a; Ernfors et al. 1990b). As with NGF and BDNF, NT-3 supports survival and neurite outgrowth from peripheral ganglia (Maisonpierre et al. 1990a; Ernfors et al. 1990b). With the exception of hippocampal dentate granule cells, NT-3 mRNA is not highly expressed in the mature CNS (Fig. 1, Allen Institute and LazC expression from the NT-3 promoter generously provided by Dr. Kevin Jones at the University of Colorado). NT-3 is present at low levels in mature animals in the indusium griseum, tenia tecta, olfactory bulb, substantia nigra, medial locus coeruleus, and scattered paraventricular (IV) brain stem nuclei (Ernfors et al. 1990b; Vigers et al. 2000). Other structures such as the retrosplenial and visual cortices with low NT-3 expression and a developmental peak were also identified in a mouse strain expressing LacZ from the NT-3 promoter (Vigers et al. 2000). LacZ expression is more sensitive than in situ hybridization, but this essentially replicated the earlier mRNA studies with respect to relative expression levels, including lack of NT-3 expression by cells in striatum and globus pallidus (Maisonpierre et al. 1990b; Friedman et al. 1991).

Adult NT-3 protein levels range from 1 to 1.5 ng/g wet weight with hippocampus, cerebellum, and piriform cortex having the highest levels and superficial and deep cortical layers containing the greatest immunoreactivity (Anderson et al. 1994). Acid treatment has been shown to increase NT-3 detection by 30–50 %, depending on the region (Okragly and Haak-Frendscho 1997). NT-3 is detected in the internal granule cell layer of the cerebellum and ELISA estimates the concentration is around 500 pg/g wet weight in mature rats (Söderström and Ebendal 1995). Levels in neonatal animals reach 9 ng/g wet weight transiently in cingulate cortex (2.5 ng/g in hippocampus) and peak expression in the hippocampus, cerebellum, piriform cortex, and entorhinal cortex occurs at birth and declines over the first 2 postnatal weeks (Ernfors et al. 1990b; Söderström and Ebendal 1995; Das et al. 2001).

NT-3 regulates the development of discrete neuronal populations during early, middle, and late periods of CNS neurogenesis and synaptogenesis. NT-3 regulates neural crest development and is expressed at the earliest steps in neuralation (Maisonpierre et al. 1990b). NT-3 has a developmental peak in mRNA expression that coincides with the arrival of perforant path afferents into the hippocampus (Maisonpierre et al. 1990b, Ernfors 1990b; Friedman et al. 1991; Lauterborn et al. 1994), and NT-3 is transiently expressed in the hilus of the hippocampus at birth (Friedman et al. 1999). Similarly, late expression of NT-3 mRNA in the visual cortex and lateral geniculate nucleus coincides with eye opening and decreases after the critical period for ocular dominance column formation to undetectable levels in the adult (Lein et al. 2000). Follow-up studies using Cre-mediated deletion in cortex establish a role for NT-3 in thalamocortical innervation (Ma et al. 2002). NT-3 expression in the thalamus is transient, and targeted deletion of NT-3 in the mouse cortex selectively perturbs the development of thalamocortical axons projecting to retrosplenial and visual cortex (Ma et al. 2002). In organotypic cultures, NT-3 induces cortical, axon fiber branching and is an attractive signal for layer 6 axons while repelling layers 2/3 axons (Castellani and Bolz 1999).

Cerebellar granule neurons switch trophism from BDNF to NT-3 after they enter the internal granule layer (Segal et al. 1992). Conditional deletion of NT-3 results in a dramatic reduction in foliation of the cerebellum with normal Purkinje cell arborization, normal EGL proliferation, and normal lamination (Bates et al. 1999). Deletion of IGF-1 receptor in the cerebellum reduces NT-3 and results in a similar foliation phenotype (de la Monte et al. 2011). In addition to scattered preand post-migratory cerebellar granule cells expressing mRNA (Rocamora et al. 1996; Ernfors et al. 1990a), NT-3 immunoreactivity is detected in cerebellar Purkinje cells, cerebellar interneurons, deep cerebellar nuclei, medial vestibular nuclei, and the inferior olive (Friedman et al. 1991; Zhou and Rush 1994). Purkinje cells have not been shown to express NT-3 mRNA (Ceccatelli et al. 1991; Ernfors et al. 1994a), suggesting that NT-3 may accumulate in Purkinje cells via retrograde or anterograde transport.

TrkC has limited distribution in the CNS of mature animals but is highly expressed early in development. TrkC binding, TrkC mRNA, and NT-3 expression are similar, with the highest receptor levels in superficial cortical layers, stratum oriens, stratum radiatum, the molecular layer of the dentate gyrus, the nucleus of the

lateral olfactory tract, entorhinal cortex, the anterior olfactory nucleus, and the anteromedial thalamic nucleus. Neocortical deep layers, striatum, amygdala, dorsal root ganglia, and the central gray of spinal cord show moderate iodinated-NT-3 binding and TrkC expression, while expression is low in the cerebellum (Altar et al. 1993, 1994; Tessarollo et al. 1993).

## 2.3.2 NT-3 and Ethanol: Animal Models

It is not surprising that few studies have examined the effect of ethanol on NT-3/TrkC expression after ethanol exposure in mature animals, given the low, limited, and discrete regional expression in the adult. Recent studies of human postmortem tissue have suggested a role for ectopic induction of NT-3 induced by oxidative stress in the etiology of autism, particularly in cerebellar dysfunction (Saidel-Sulkowska et al. 2011), but studies on NT-3 in developing and adult animals have been equivocal. NT-3 administered into the ventricles did not rescue vasoactive intestinal peptideimmunoreactive neurons in the suprachiasmic nucleus of withdrawn animals (Paula-Barbosa et al. 2003). Similarly, Baek and colleagues (1996) did no observe any changes in hippocampal NT-3 mRNA normalized to cyclophilin A after 28 weeks of constant ethanol exposure via liquid diet. Miller (2004) used a repeated episodic vapor exposure model, while Miller and Mooney (2004) exposed rats chronically to ethanol via liquid diet for 8 or 24 weeks and measured NT-3 levels by ELISA in parietal and entorhinal cortices, hippocampus, septum, and basal forebrain. Episodic exposure increased hippocampal NT-3 levels (from  $\sim 100$  to 155–195 pg/mg protein) and reduced in basal forebrain NT-3 (from  $\sim$ 500 pg/mg protein to  $\sim$ 200 pg/mg). Chronic exposure increased NT-3 in entorhinal cortex and septum at both 8 and 24 weeks but transiently increased levels in basal nuclei and hippocampus. Both Miller and Mooney (2004) and Miller (2004) report the highest NT-3 content in the septum (600–700 pg/mg protein) with hippocampal levels ranging from 100 to 150 pg/mg protein and entorhinal cortex slightly higher at  $\sim$ 200 pg/mg protein. These data are given normalized to total protein but assuming 10 % protein, a value of 100 pg/mg is roughly 10 ng/g wet weight and regions such as septum contain 45+ ng/g wet weight. It should be noted that these concentrations are higher and not in the regional proportions previously observed in the adult, even upon acid treatment (see above) but may be the result of different sample preparation methods or, alternatively, significant transport.

When measured in homogenates after a variety of exposure paradigms, ethanol exposure either prenatally or postnatally changed NT-3 mRNA (Fattori et al. 2008; Light et al. 2001; Tsuji et al. 2008) and protein levels measured by ELISA (Heaton et al. 2000c). Heaton and colleagues (2000c) showed a trend toward increased "cortex/striatum" NT-3 on PN10 with PN4-10 vapor exposure (~90 pg/g wet weight to ~140 pg/g wet weight) in pups, but this did not reach statistical significance and NT-3 levels did not differ in other regions examined. When the telencephalon was further dissected (Heaton et al. 2003b), exposure to ethanol at PN14, but not the vulnerable PN3 exposure, increased NT-3 from ~8 ng/g to ~14 ng/g of striatal wet weight in rats at 24 h after exposure; NT-3 levels did not vary between

PN3 and PN14 (Heaton et al. 2003b). These levels are perplexing, given the variable concentrations observed between experiments and the lack of a developmental decrease in NT-3 postnatally (i.e., Das et al. 2001). NT-3 mRNA and NT-3 promoter-driven LacZ is not detected in the striatum in developing or mature animals (Maisonpierre et al. 1990b; Vigers et al. 2000; Soderstrom and Ebendal 1995; Das et al. 2001; Ernfors 1990b; Friedman et al. 1991; Lauterborn et al. 1994). Moreover, mRNA and LacZ expression also indicate low level of expression from the resident cells in the septum and basal forebrain. Therefore, the source of striatal NT-3 is unclear in these experiments as well. As discussed above, NT-3 is transiently produced in many structures including the thalamus, visual, cingulate, and retrosplenial cortex. NT-3 also regulates thalamocortical axon development during the critical stage with a peak in mRNA expression at PN8 (Ma et al. 2002). Again, NT-3 protein measured in striatum could be in retrograde transit through the striatum since the PN3 and PN14 time points in Heaton (2003b) flank the peak at PN5-8 but levels have not been shown to vary by orders of magnitude within a few days developmentally.

When the cerebellum is subdivided or immunoreactivity evaluated in situ, NT-3 and TrkC are sensitive to ethanol exposure in selected regions. TrkC immunoreactivity is reduced when examined specifically on Purkinje cells in pups exposed to ethanol during the vulnerable PN4-6 window (Light et al. 2002), but global mRNA for TrkC, likely representing mRNA from mature granule cells as well, was unchanged (Light et al. 2001). When the vermis was selectively assayed by ELISA, Parks and colleagues (2008) found increased cerebellar NT-3 and an increase in hippocampal NT-3 in mature animals after prenatal exposure, which could reflect compensation within the system. The high dose (6 g/kg/day by gavage) increased NT-3 in both regions while only the vermis was sensitive to the lower dose (4 g/kg/day). In addition, NT-3 levels in the vermis were higher in animals exposed to an enriched environment after prenatal alcohol exposure at both doses, again suggestive of a role in plasticity. The NT-3 concentrations measured were again higher than earlier reports in normal rats, but the relative levels, with the hippocampus having the highest NT-3 content, followed by cerebellum, were consistent.

The discrepancy between these studies and the classical descriptions is unclear as is the source of the high NT-3 levels measured in tissues not shown previously to have high levels of NT-3, suggesting the source of the NT-3 may be external to the structure. Acid treatment has been shown to increase NT-3 detection, but this should be uniform and not change the relative levels from region to region. Neutralization can also cause the reformation of NGF protein complexes. The samples in Miller and Mooney (2004) and Miller (2004) were fractionated by centrifugation, which may have concentrated specific pools of NT-3, and acid treated prior to assay; the samples in the Heaton studies were also acid treated. Parks and colleagues report higher levels than previously observed, but ratios consistent with the classical studies and those using in acid treated/neutralized samples in normal rodents.

### 2.3.3 NT-3 and Ethanol: In Vitro Models

The role of NT-3 in development is similarly time- and population-selective, and the effect of NT-3 on ethanol toxicity is similarly selective. Seabold and colleagues (1998) did not observe a protective effect of NT-3 on differentiated cortical neurons treated in vitro with a high (400 mg/dL) concentration of ethanol. Earlier in development, exposure of embryo cultures to ethanol for 3–10 h in vitro reduced proliferation of cephalic neural crest cells that was prevented by the addition of NT-3 to the medium (Jaurena et al. 2011). These studies are consistent with the limited cortical responsiveness to NT-3 and the early developmental peak in expression (see Sect. 2.3). Recent studies indicate that NT-3 may have a larger role in cerebellar development than previously appreciated. Cerebellar foliation is impaired when the insulin or IGF-1 system is perturbed in the cerebellum by siRNA; NT-3 levels are also reduced (de la Monte et al. 2011). This treatment reduces cerebellar foliation, suggesting role for NT-3 in this process, and developmental ethanol exposure similarly impairs cerebellar foliation (Cragg and Phillips 1985; Sakata-Haga et al. 2001).

# 3 Nonclassical Neurotrophic Factors: IGF-1, GDNF, and PACAP

## 3.1 The Insulin Family: IGF and ALK

IGF-1 is known to be involved in ethanol toxicity both centrally and peripherally. IGF-1 is produced in the liver in response to growth hormone secretion from the pituitary and thereby links IGF-1 levels to ethanol-induced growth restriction. Circulating IGF-1 levels in adults may also be involved in ethanol-induced vascular damage due to a lack of IGF-1-mediated repair, either directly by inhibiting signaling or secondary to hepatotoxicity, thereby compromising the blood-brain barrier. IGF-1 is bound in vivo to IGF Binding Proteins (IGFBPs) that further regulate bioavailability. Acute ethanol exposure results in a dramatic increase in IGFBP-1 and a decline in IGF-1 in healthy adults, while chronic ethanol exposure decreases IGF-1 bioavailability and growth hormone levels in patients (Passilta et al. 1999; Rojdmark et al. 2000; Rojdmark and Brismar 2001). After 4 days of withdrawal, IGF levels increase and IGFBP-1 levels decrease (Passilta et al. 1998). IGF-I and IGF-II are both upregulated in children exposed prenatally to ethanol, indicative of compensatory induction and a potential biomarker for FAS/ARND (Aros et al. 2011). The significance of alterations in peripheral IGF-1 to brain pathology is uncertain, but IGF-1 crosses the blood-brain barrier (Pan and Kastin 2000) and transport is activity dependent (Nishijima et al. 2010). Moreover, reduced IGF-1 levels are associated with depressive-like behavior (Mitschelen et al. 2011). Circulating IGF-1 levels are sensitive to hepatic function, thereby providing a link between hepatotoxicity and neuronal toxicity and repair processes. The GH-IGF-1 axis, neural activity, vascular integrity, and hepatic function are intimately linked, particularly during development.

IGF-1 is active in the early stages of neurogenesis, regulates stem cell proliferation, and enhances survival after ethanol exposure (Tateno et al. 2004). Ethanol blocks IGF-1-stimulated proliferation in vitro (Resnicoff et al. 1996) and ERK activation (Hallak et al. 2001) but, while trophic, does not significantly protect cerebellar granule cells from ethanol toxicity (Bonthius et al. 2003). Interestingly, in vivo siRNA-mediated reduction in insulin or IGF-1 receptor signaling produces cerebellar malformations that resemble fetal alcohol syndrome (de la Monte et al. 2011). McClure and colleagues (2011) used Drosophila to model fetal alcohol syndrome and implicate *Drosophila* insulin-like peptide and reduced insulin receptor expression in the teratogenic effects of ethanol. Another related molecule, ALK, was also recently suggested to be ethanol-sensitive in fly and mouse experiments (Lasek et al. 2011a, b). ALK is proapoptotic in the absence of ligand, but signals survival in presence of ligand (Yanagisawa et al. 2010). ALK promotes sensitization and cocaine reward (Lasek et al. 2011a) in flies and is involved in ethanol tolerance and reward in mice (Lasek et al. 2011a, b). ALK knockout mice show enhanced loss of righting reflex in response to high doses of ethanol, and this is not explained by reduced ethanol metabolism. ALK expression in striatum is negatively correlated with ethanol sensitivity. While knockout mice were more sensitive to the sedating effects, they were less sensitive to ataxia. These data implicate multiple intermediates in the IGF pathway as potential modulators of ethanol sensitivity and toxicity.

# 3.2 GDNF

GDNF is also a target derived factor that was originally described as a trophic factor for dopaminergic, sensory, and motor neurons (reviewed by Pascual 2011). GDNF is a member of a larger family that includes Artemin, Neurturin, and Persephin. While trophic for dopaminergic neurons, genetic ablation of GDNF does not affect genesis or survival of dopaminergic neurons (Sanchez et al. 1996; Jain et al. 2006), but Ret deletion cause late degeneration of the substantia nigra pars compacta (Kramer et al. 2007). GDNF is also produced in peripheral tissues, with high levels detected in kidney, ovary, testes, stomach, skin, and lung (Trupp et al. 1995). Like neurotrophins, GDNF signals through tyrosine kinase coupled pathway. GFR-alpha 1 binds GDNF and recruits the Ret tyrosine kinase; neurturin preferentially activates alpha 2, and artemin preferentially activates alpha 3 (reviewed by Mason 2000). Ret activation leads to ERK, Akt, phospholipase C, and Src activation to promote survival and differentiation. GDNF is also a potent regulator of antiapoptosis pathways and NFkB. As with classical neurotrophins, alternative signaling pathways also exist for GDNF. GDNF can signal through NCAM to activate Fyn and focal adhesion kinase which mediate neurite outgrowth and survival (Sariola and Saarma 2003). Homophilic transsynaptic binding of GFR-alpha has also been described during hippocampal synaptogenesis (Ledda 2007).

GDNF is expressed at high levels in the striatal fast spiking interneurons, olfactory bulb, ventral mesencephalon, and septum with lower expression in the cerebellum and colliculi (see Fig. 1 and Hidalgo-Figueroa et al. 2012;

Pochon et al. 1997; Golden et al. 1999 for detailed anatomy). Receptors for GDNF are more widely expressed with Ret mRNA detected at high levels in diencephalic structures, cerebellum, brain stem, and olfactory bulb. Ret expression is low in the hippocampus, cortex, and striatum. GFR-alpha 1 is expressed at high levels in the septum, hippocampus, and colliculi, but at lower levels in the olfactory bulb, diencephalon, and brainstem. GFR-alpha 2 is high in cortex, hypothalamus, and colliculi, with low expression in the cerebellum and brainstem. GFR-alpha 3 expression is limited to the developing CNS and cerebellum (Masure et al. 1998) but is expressed in peripheral tissues such as heart, lung, and trigeminal ganglia. Interestingly, the striatal FSIs, not medium spiny neurons, produce GDNF while the receptors are produced by, and GDNF is accumulated in, mesencephalic dopamine neurons (Hidalgo-Figueroa et al. 2012; Barroso-Chinea et al. 2005). As with other neurotrophins, the differential distribution of GDNF and its receptors is consistent with target-derived factors and the consumer-producer model of neurotrophin action.

As with BDNF, GDNF was also identified for having the ability to reduce alcohol consumption in rodents (Carnicella et al. 2008, 2009a, b). The locus for this effect is the VTA since infusion of GDNF shRNA into the VTA blocks the effect (Barak et al. 2011) so it does not appear to involve fast spiking interneuron in the striatum, which also produce GDNF. Examination of the effects of ethanol on GDNF-mediated survival is limited. Fiore and colleagues (2009b) observed a decrease in cortical GDNF after a diet of chronic red wine (but not ethanol). Hippocampal GDNF does not appear sensitive to ethanol in mature animals (Okamoto et al. 2006).

Developmental studies in vivo indicate that GDNF protects developing chick spinal motor neurons from ethanol, but ethanol exposure did not reduce muscle GDNF levels in rats (Bradley et al. 1999; Barrow Heaton et al. 1999). However, Wentzel and Ericksson (2009) found a decrease in GDNF in trunk, but not cranial, neural crest–derived cells from the E10 rat embryo. Interestingly, this paper also reported a decrease in GAPDH mRNA in cranial neural crest cells exposed to ethanol. McAlhany and colleagues (1999) also found a reduction in GDNF protein, but not mRNA, with developmental ethanol exposure but a paradoxical increase in GDNF signaling to Shc. Neonatal brain growth restriction in response to high dose (5 g/kg gavage) ethanol during the postnatal brain growth spurt (PN6-8) is similarly correlated with a transient reduction in GDNF (Tsuji et al. 2008). (Exposure on PN 2–4 or PN 13–15 also reduced BDNF levels, but this did not correlate with microcephaly, as discussed above.) Interestingly, the ERK pathway was similarly sensitive regardless of age but inhibition of the Akt pathway was only observed with PN6-8 exposure, correlating with the temporal decrease in GDNF in these experiments.

In vitro, GDNF reverses Purkinje cell ethanol toxicity in cerebellar explants (McAlhany et al. 1997, 1999) and ethanol blocks GDNF-stimulated process outgrowth in granule cells (Chen and Charness 2012). GDNF reversed ethanol-induced JNK activation in SK-N-SH neuroblastoma cells and associated cell death (McAlhany et al. 2000). Interestingly, these cells underwent DNA fragmentation at 34 and 68 mM ethanol but displayed signs of necrosis at concentrations above 100 mM. GDNF can also protect glial-type transformed cells from ethanol toxicity in vitro (Villegas et al. 2006). In contrast, Mitchell and colleagues (1999) found no neuroprotective effects of GDNF against cell death in hippocampal cultures, induced by high concentrations (>100 mM, 0.4–1.6 g/dl) of ethanol combined with hypoglycemia and hypoxia.

## 3.3 PACAP

PACAP is a member of the vasoactive intestinal peptide/glucagon/growth hormone releasing hormone superfamily and is active in stress, anxiety, consummatory behavior, metabolism, and thermoregulation (Adams et al. 2008). PACAP receptors exists as splice variants that stimulate adenylate cyclase and phosphoinositol hydrolysis with different kinetics (Spengler et al. 1993). PACAP also signals through Ca<sup>++</sup>, PCK; Src family kinases, and by transactivation of Trks through the PAC1 receptor, with Gq-coupled pathways predominating in the hippocampus (Macdonald et al. 2005; Rajagopal et al. 2004; Shi et al. 2010). PACAP may also regulate BDNF levels through receptor-coupled activation of RACK1 through cAMP (Yaka et al. 2003a) placing BDNF downstream of PACAP. PACAPdeficient mice display hyperlocomotion, decreased body mass, novelty-seeking behavior, jumping, impaired hippocampal associative learning, and mossy fiber LTP. PACAP in the arcuate nucleus of the hypothalamus stimulates NPY neurons to regulate carbohydrate intake (Nakata et al. 2004) but inhibits food intake via the POMC arcuate neurons (Mounien et al. 2009). Peripherally, PACAP is coupled to catecholamine secretion from adrenal chromaffin, providing a peripheral link to the stress response (Kuri et al. 2009).

The initial evidence for involvement of PACAP in ethanol sensitivity came from *Drosophila* where an allele of the *Drosophila* PACAP homologue amnesiac called cheapdate was identified as a gene regulating ethanol sensitivity (Moore et al. 1998). Rutabaga, another cAMP pathway mutant, is required in mushroom bodies for ethanol consumption in *Drosophila* (Xu et al. 2012). As with flies, PACAP knockout mice are less sensitive to the hypothermic and hypnotic effects of ethanol (Tanaka et al. 2004). Cerebellar granule cells from PACAP knockout mice are more sensitive to ethanol and peroxide toxicity, while addition of PACAP or dibutryl cyclic AMP reversed this effect (Vaudry et al. 2005). PACAP administered to neonatal rats mitigated the ethanol-induced loss of righting and apoptosis, possibly through caspase 3 and cJun inhibition (Botia et al. 2011). As with BDNF, PACAP also protected cerebellar granule cells from ethanol toxicity by the activation of PI3K (Bhave and Hoffman 2004).

# 3.4 Nonconventional Neurotrophic Factors in Patients with Alcohol Dependence

To date only one study has examined peripheral GDNF levels in alcohol-dependent patients. Heberlein and colleagues (2010) found GDNF levels significantly

depressed in the serum of alcohol-dependent patients which remain decreased for up to 2 weeks after withdrawal. Interestingly, serum levels were also inversely related to subjective assessment of tolerance. Those with high serum GDNF levels reported lower tolerance and may therefore be more likely to limit their alcohol intake. This is consistent with observations in rodents, suggesting that GDNF negatively regulates ethanol consumption (Carnicella et al. 2009a, b).

# 4 Neurotrophic Therapies in Alcoholism, FAS, and ARND

Neurotrophins-based therapies for both addiction and repair of ethanol-induced brain damage may be on the horizon. Small interfering RNA and other expressionbased strategies have been used experimentally to dissect the pathways involved in ethanol self-administration in rodents, but this strategy will not be immediately useful for treating patients. Direct activation of Trks with ICV neurotrophin infusion has been used experimentally. For example, neurotrophin infusion can maintain ethanol tolerance (Szabò and Hoffman 1995) and infusion of NGF can restore cholinergic function in the mature brain after chronic ethanol exposure in rodents (Cadete-Leite et al. 2003; Lukoyanov et al. 2003). Unfortunately, ICV infusion of NGF is also not feasible for human treatment, but intranasal delivery of IGF-1 is being investigated for ameliorating ischemia and hypoxia-induced damage in developing mice (Lin et al. 2009, 2011) and in adult stroke models (Kooijman et al. 2009). Intranasal IGF-1 also protects Purkinje cells in spinocerebellar ataxis type 1 and improves roto-rod function (Vig et al. 2006). Given the trophic role of IGF-1 in the CNS, IGF-1 might also be useful in FAS/ARND and it readily crosses the blood-brain barrier. While not protective against the insult, IGF-1 may enhance recovery in the child.

Neurotrophic peptides and peptidomimetics are also being developed, which may reverse ethanol-induced neurodegeneration and withdrawal-associated anxiety. Cyclotraxin-B was recently isolated from a peptidomimetic screen as a TrkB allosteric antagonist that reduces anxiety-like behavior (Cazorla et al. 2010, 2011). Such molecules may reduce anxiety-associated drinking and withdrawal hyperexcitability but may also pose a toxicity risk. Conversely, mimetic and small molecule agonists of TrkA and TrkB reverse cell death and enhance recovery in experimental models (Scarpi et al. 2012; Massa et al. 2010; Jang et al. 2010) and may therefore be useful for treating ethanol-induced neurodegeneration and cognitive dysfunction. Activity-Dependent Neurotrophic Factor and related peptides enhance survival in moderate (Zhou et al. 2008) and extreme (Incerti et al. 2010) fetal alcohol models. Interestingly, the ADNF peptides SAL and NAP reversed the deficits observed after a single exposure on day 8 of gestation to a dose of ethanol that produces BACs in excess of 500 mg/dL. This protective effect was attributed to normalization of ethanol-induced upregulation of BDNF at E9 and E18. Thus, NAP and SAL can influence events prior to the onset of CNS neurogenesis and normalize BDNF expression without off target teratogenesis; however, recent studies have excluded a direct role for BDNF in NAP effects on axon formation (Chen and Charness 2012). The efficacy of NAP and related peptides may therefore be related to cytoskeletal interactions (Gozes 2011), or inhibition of apoptosis (Zemlyak et al. 2009) and not a direct interaction with BDNF signaling.

pharmacological strategies that indirectly Small molecule modulate neurotrophin levels have also been proposed and tested in rodent models. Antidepressants increase BDNF in the hippocampus and can reverse ethanol-induced reductions in BDNF in the hippocampus (Hauser et al. 2011), which may increase recovery after long-term ethanol exposure. The anti-addiction compounds Ibogaine and noribogaine may also interact with GDNF to reduce ethanol drinking (He et al. 2005; Carnicella et al. 2010). Neurotrophin-based therapeutics need not involve administration of a drug, however. Exercise in mature animals can induce hippocampal neurogenesis and increase BDNF levels, which ameliorates many of the effects of prenatal ethanol exposure (Boehme et al. 2011). Antioxidants provide neuroprotection in fetal alcohol models, in part through modulation of neurotrophins (reviewed by Brocardo et al. 2011). Dietary supplementation with omega-3 fatty acids also reverses behavioral deficits in fetal alcohol models and increases BDNF, possibly through antioxidant modulation (Patten et al. 2012). Similarly, choline supplementation modulates neurotrophins and choline is being examined in preclinical models as a treatment for alcohol-related birth defects (Ryan et al. 2008; Monk et al. 2012). These interventions, while subtle, may increase neurogenesis and plasticity, ultimately involving regulation of neurotrophins. The strategies vary but all ultimately increase neurotrophic activity to influence survival and plasticity.

## 5 Conclusion

Alterations in neurotrophins and neurotrophin signaling mediate the acute toxic effects, plasticity that occurs with pharmacodynamic tolerance, plasticity that occurs with addiction, and repair that occurs after withdrawal. The temporal and cell specific activation and inhibition of neurotrophins and their associated pathways have not been conclusively determined, and the data on ethanol and neurotrophic factors summarized in this chapter clearly do not outline a cohesive story. Neurotrophin levels can increase, decrease, or not change depending on the brain region, time of exposure, and presence or absence of drug upon analysis. Neurotrophin signaling is dependent upon concentration, with full-length receptors desensitizing, high concentrations activating p75, and related receptors being nonspecifically activated at higher concentrations. In addition, neurotrophin levels and signaling to not occur in isolation; therefore, it is important to consider the physiological changes that result from neurotrophin action. For example, an increase in glutamatergic synaptic activity during withdrawal would be expected to increase BDNF but the place of BDNF in this sequence is not well defined in most studies. Does BDNF drive these changes or, more likely, does BDNF increase as a result of excess NMDA receptor activation? The sequence is different early in exposure and with low ethanol concentrations where ethanol can reduce signaling in BDNF independent of effects on ion channels (Kolb et al. 2005; Ohrtman et al. 2006). In this case, neurotrophin signaling may be directly involved in the regulation of subsequent events and not vice versa. When concentrations of ethanol become too high, none of these adaptive responses are likely to occur, however, as is observed with suprapharmacological concentrations. More subtle effects are likely to contribute to ethanol-induced teratogenesis and chronic toxicity.

The expression data for receptors and neurotrophin levels measured after ethanol exposure also frequently do not agree. It is possible that ethanol induces ectopic expression in specific cell types, but this has not been specifically addressed since the control cells in these regions also frequently show expression of neurotrophins they are not believed to express. This may be partially due to the fixation procedures used to detect immunoreactivity in situ since neurotrophins are difficult to detect with immunohistochemistry. In situ hybridization and protein detection with and without colchicine treatment and global mapping may also help resolve some of these discrepancies and locate the "producer" neurons where neurotrophins are sensitive to ethanol treatment. The availability of genetically tagged mice will also aid in dissecting these issues anatomically, functionally, and allow for separation of specific populations by cell sorting. In addition, Cre-recombinase-mediated deletion in specific cell types, for example, BDNF deletion in cortex versus medium spiny neurons, could be combined with self-administration behavioral experiments to define the loci for specific effects. The pharmacological tools described in the previous section will aid in dissection of the actions of neurotrophins in ethanol toxicity, tolerance, addiction, withdrawal, and the development of FAS/ARND.

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# Neurotrophic Factors and NeuroAIDS: A Lesson from Brain-Derived Neurotrophic Factor

Alessia Bachis, Valeriya Avdoshina, Seung T. Lim, and Italo Mocchetti

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#### Abstract

Human immunodeficiency virus type 1 (HIV) causes mild or severe neurological problems, termed HIV-associated neurocognitive disorder (HAND), even when HIV patients receive antiretroviral therapy. Thus, novel adjunctive therapies are necessary to reduce or abolish the neurotoxic effect of HIV. However, new therapies require a better understanding of the molecular and cellular mechanisms of HIV-induced neurotoxicity. HAND subjects are characterized by being profoundly depressed, and they experience deficits in memory, learning, and movements. These deficits resemble those occurring in premature brain aging.

A. Bachis (🖂) • V. Avdoshina • S.T. Lim • I. Mocchetti

Department of Neuroscience, Georgetown University Medical Center, Washington, DC, USA e-mail: bachisa@georgetown.edu; va44@georgetown.edu; ls379@georgetown.edu; moccheti@georgetown.edu

Thus, it appears that HIV diminishes neuronal survival, along with reduced neuronal connections. These two phenomena should not occur in the adult brain when synaptic plasticity is promoted by neurotrophic factors. In particular, synaptic plasticity is enhanced by brain-derived neurotrophic factor (BDNF), a potent neurotrophic factor that is present in abundance in the adult synapses. This chapter will outline experimental evidence as well as present emerging concepts for the use of BDNF as an adjunct therapy to prevent HIV-mediated neuronal degeneration and restore the loss of synaptic connections.

#### Keywords

Acquired immunodeficiency syndrome • Adeno associated viruses • Apoptosis • Basal ganglia • BDNF synthesis • Brain-derived neurotrophic factor • Caspase-3 • CCR5 • CXCR4 • Dendrite branching • Dendritic • Excitotoxicity • Frontal cortex • gp120 • HIV-associated dementia • HIV-associated neurocognitive disorders • HIV encephalitis • Human immunodeficiency virus type-1 • Microglia • Nerve growth factor • Neurotrophic factors • Neurotrophins • Nigrostriatal dopamine neurons • p75NTR • Proneurotrophins • Simplification • Synaptodendritic pathology • TrkB • Viral protein

# 1 Introduction

*Human immunodeficiency virus type 1* (HIV) which depletes the immune system causing *acquired immunodeficiency syndrome* (AIDS) also infects the central nervous system (CNS) and promotes neurological problems in more than 50 % of patients not receiving any form of antiretroviral therapy. Symptoms include profound motor and behavioral/psychosocial abnormalities that disrupt work or other activities of daily living. Abnormalities that can be manifested as attention deficit can also be seen in children.

The introduction of combined antiretroviral therapy (cART) has improved immune recovery and decreased the gravity of neurological signs and the more severe form of cognitive impairment (Sacktor et al. 1999) or HIV-associated dementia (HAD). By controlling the viral load, cART allows an individual with HIV to typically live longer with milder medical symptoms. Indeed, recent estimates of AIDS subjects after the advent of cART place HAD in the 1-2 % of the HIV cohort (Woods et al. 2009). Yet, cART has not eliminated mild neurocognitive deficits and asymptomatic neurocognitive impairments (McArthur et al. 2003). These deficits are referred to as HIV-associated neurocognitive disorders (HAND). Current estimates find that nearly 50 % of HIV patients in the United States demonstrate neuropsychological testing performance that is below expectations compared to age-, education-, gender-, and ethnicity-matched reference groups. The reason why HAND persists among treated individuals is still under debate. Some of the drugs used in cART may not reduce HIV infection in the CNS because they poorly penetrate the blood-brain barrier (BBB) or because the virus develops drug resistance. It is noteworthy that neurologically impaired individuals have a higher viral load in the cerebrospinal fluid (CSF) than in plasma. Indeed, clinical evidence has shown that the CNS of these individuals is an HIV sanctuary site (Masliah et al. 2000) unless they initiate a highly CNS-penetrating antiretroviral regimen (Letendre et al. 2004).

HIV does not infect neurons; nevertheless, postmortem brains of HAD subjects have shown neuronal loss accompanied by synaptic simplifications (Everall et al. 2005). Thus, our great hope lies in our ability to develop alternative adjunct therapies that can provide a more efficient treatment of HIV-mediated neuronal degeneration. The search for pharmacological compounds that prevent HIV neurotoxicity must rely on the general concept of the so-called biological therapies, which are defined by two principal features: they require a better understanding of the molecular and cellular mechanisms of HIV neurotoxicity, and, ideally, they use physiological compounds utilized by the CNS to prevent neuronal injury. These compounds include *neurotrophic factors*, which are naturally occurring diffusible polypeptides that promote survival of a variety of CNS cells and are equally essential for inducing differentiation of surviving neurons into their mature phenotypes (Reichardt 2001). This chapter will introduce new emerging concepts and principles in the use of neurotrophic factors as an adjunct therapy to prevent HIV-mediated neuronal degeneration and restore the loss of synaptic connections.

## 2 Brain-Derived Neurotrophic Factor: The Most Abundant Neurotrophin in the Adult Brain

#### 2.1 Neurotrophins

HAND pathology includes loss of both synaptic connections and neuronal differentiation (Ellis et al. 2007). In addition, HIV promotes neuronal apoptosis, especially in children (Gelbard and Epstein 1995). Thus, this disease is an excellent target for a neurotrophic factors-based therapy. In fact, neurotrophic factors have been shown to reduce synaptic and axonal degeneration mediated by a number of neurotoxins and to interfere with the fundamental mechanism of apoptotic cell death in numerous conditions (Reichardt 2001). One of the most potent neurotrophic factors is *brain-derived neurotrophic factor* (BDNF) (Barde et al. 1982), the most abundant member of the neurotrophin family of trophic factors in the adult CNS that also includes *nerve growth factor* (NGF) (Levi-Montalcini 1987), neurotrophin-3, and neurotrophin-4 (Maisonpierre et al. 1990).

BDNF exerts multiple neurotrophic activities during development as well as in the adult CNS. After its initial discovery by Barde et al. (1982), interest in BDNF grew due to its potent effect on cortical (Cabelli et al. 1995) and subcortical plasticity (Lyons et al. 1999; Mamounas et al. 1995). Neurotrophic effects include modulation of dendritic branching and spines in the cortex and long-term potentiation in the hippocampus (Patterson et al. 1996; Zakharenko et al. 2003). Through these properties, BDNF plays a critical role in learning and memory and preservation of cortical circuits. Conversely, a reduction of BDNF secretion/activity is responsible for the loss of cortical and hippocampal synapses and fear learning. This has been demonstrated not only in animals but also in humans. In fact, impaired learning and memory have been observed in both human subjects (Egan et al. 2003) and mice (Soliman et al. 2010) in which the regulated BDNF secretion is reduced due to a single nucleotide polymorphism in the BDNF gene that encodes a valine (Val) to methionine (Met) substitution at codon 66 (Val66Met). In addition, correlative studies of human chronic neurodegenerative diseases characterized by reduced BDNF levels/activity have confirmed the role of BDNF in adult plasticity. For instance, a deficiency in BDNF synthesis has been described in postmortem brains (Phillips et al. 1991) or cerebrospinal fluid (Laske et al. 2007) of patients with Alzheimer's disease (AD), which is categorized as a disease with reduced number of neurons and connections in the cortex and hippocampus as well as impaired short-term memory (Scheff and Price 2006).

BDNF is also decreased in nigrostriatal dopamine neurons in Parkinson's disease (PD) (Howells et al. 2000; Nagatsu et al. 2000), cortical neurons in Huntington's disease (HD) (Zuccato et al. 2001), and schizophrenia (Durany and Thome 2004). Nevertheless, results obtained from these studies may reflect secondary effects of the main pathological process rather than primary pathological events. This limitation is especially problematic for the *neurotrophins* whose expression and availability is regulated by synaptic activity (Ghosh et al. 1994). However, BDNF treatment has the potential to reduce or abolish neuronal death in several brain areas. Indeed, early experimental studies in animal models have shown a remarkable neuroprotective effect of BDNF against various neurotoxins that mimic human chronic neurodegenerative diseases such as PD and HD (Alberch et al. 2002; Frim et al. 1994; Hyman et al. 1991). Similar neuroprotection has been observed in rodents and nonhuman primates after the lesion of the perforant path, an animal model of AD (Nagahara et al. 2009). These and other experimental evidence have prompted the suggestion that BDNF could be a potential therapeutic agent for the treatment of these diseases. Most importantly, these observations suggest that an environment characterized by lower BDNF levels or activity (and other neurotrophic factors) could be a common risk factor for the development of neurological diseases. This issue rarely receives the attention it deserves, a problem that may have a negative impact on the success of future clinical trials.

# 2.2 Neurotrophin Receptors: Dual Mechanism of Action

The biological activity of the neurotrophins begins when they bind to a receptor complex composed of two different receptors. The first neurotrophin receptor that was characterized was a member of the tumor necrosis factor receptor family that was named p75 NGF (Johnson et al. 1986; Radeke et al. 1987); however, p75 binds to all neurotrophins with a similar affinity (Rodriguez-Tebar et al. 1990) and was therefore named *p75NTR*. The other component of the neurotrophin receptor complex is the proto-oncogene Trk. This is a receptor tyrosine kinase which, like other tyrosine kinase receptors, is activated by ligand-induced formation of non-covalently

associated receptor dimers (Kaplan et al. 1991). There are three structurally related Trks and each neurotrophin binds selectively to Trk: BDNF binds to TrkB, NGF to TrkA, and NT-3 to TrkC (Chao and Hempstead 1995); however, at high concentrations BDNF can also bind to TrkC (Klein et al. 1991). Both Trk and p75NTR are necessary to confer high-affinity binding to the neurotrophins and to influence their biological activities (Brennan et al. 1999; Hempstead et al. 1991). When activation of p75NTR occurs without a concomitant activation of Trk, p75NTR promotes death of oligodendrocytes (Gu et al. 1999) as well as axonal degeneration in the peripheral (Kenchappa et al. 2006) and CNS (Park et al. 2010). In addition to Trk, p75NTR can also form a complex with the so-called truncated Trk (or Trk.T1), an isoform of Trk generated from an alternative splicing which does not contain the tyrosine kinase domain. In certain neuronal populations, TrkB.T1 or TrkC.T1 can function as inhibitors of full-length Trk through a dominant-negative mechanism. Nevertheless, Trk isoforms and p75NTR also exhibit some neurotrophic properties such as neuronal crest proliferation and differentiation (Hapner et al. 1998) and regulation of neuronal branching as well as BDNF signaling (Carim-Todd et al. 2009).

p75NTR can also bind the larger precursor protein for the neurotrophins or proneurotrophins including proBDNF (Hempstead 2002). Proneurotrophins are abundant in the mature brain (Fahnestock et al. 2001) and can be released from neurons (Yang et al. 2009). Proneurotrophins can be cleaved in the endoplasmic reticulum by the proconvertase furin (Seidah et al. 1996) or extracellularly by proteases such as plasmin and matrix metalloproteases to be converted to mature neurotrophins (Mowla et al. 1999). Conversion of proneurotrophin to mature neurotrophin is an important process for synaptic plasticity. In fact, when processing is impaired, proNGF or proBDNF can accumulate outside neurons and induce neuronal loss by binding to p75NTR (Harrington et al. 2004; Teng et al. 2005). The mechanisms of p75NTR-mediated cell death include several "toxic" signaling pathways that are unrelated to tyrosine kinase activation but that produce ceramide (Dobrowsky et al. 1994) and activate nuclear factor kB and Jun kinase (Bhakar et al. 2003; Carter et al. 1996). Given the fact that at high concentrations neurotrophins can indiscriminately activate p75NTR, caution must be exerted in delivering neurotrophins into the mature injured brain where there is a possibility of an upregulation of p75NTR without a concomitant increase in Trk (Beattie et al. 2002; Brunello et al. 1990; Volosin et al. 2006).

# 3 BDNF and HIV Dementia: Basic Tenets

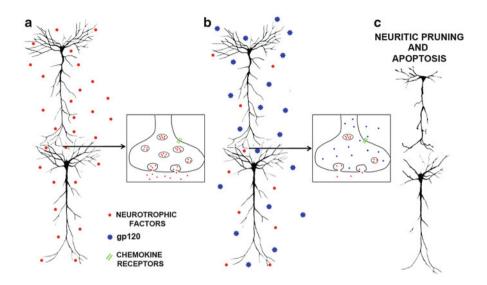
Subjects with the most severe form of HAND or HAD exhibit depression and mania (Grant 1990; Pumpradit et al. 2010), combined with all degrees of cognitive and motor impairments. In fact, the clinical manifestations of HAD include tremor, gait ataxia, loss of fine motor movement, mental slowing, forgetfulness, poor concentration, and behavioral abnormalities (McArthur et al. 2010). Neuroimaging studies of HAD patients have revealed generalized white matter reduction, with additional grey matter loss particularly in the basal ganglia and posterior cortex

(Aylward et al. 1995; Dal Pan et al. 1992). Neuronal loss has been confirmed in postmortem brains in the *basal ganglia* and other regions of the brain including the hippocampus and *frontal cortex* (Davies et al. 1998; Everall et al. 1993). Pathological alterations of the basal ganglia in HAD include neuronal loss in the putamen (Everall et al. 2005) and the globus pallidus (Fox et al. 1997), degeneration of nigrostriatal dopamine (DA) neurons (Itoh et al. 2000; Reyes et al. 1991), and dysfunctional DAergic transport (Wang et al. 2004). Thus, it is not surprising to note clinical features in HAD resembling those found in PD, such as the postural instability, involuntary movements, bradykinesia, and impairment in fine motor skills (Berger and Nath 1997; Koutsilieri et al. 2002; McArthur 2004).

HAD also exhibits a number of striking similarities with AD in terms of clinical manifestations and neuropathology. Indeed, cognitive impairments (moderate to severe) in HAD are similar to those observed in AD. Higher cognitive functions depend on a highly complex synaptodendritic network in the frontal cortex, and damage to this network results in abnormal output, measured as deficiencies in cognitive skills and behavior. Granted that memory and attention alterations could be also due to the high prevalence of CNS comorbidities in HIV populations (e.g., drug abuse), it is important to keep in mind that these cognitive alterations correlate with the loss of synaptodendritic networks in the cortex and hippocampus. Moreover, neuroimaging studies have revealed microstructural abnormalities in the cerebral white matter, supporting the theory of *synaptodendritic pathology* as the main cause of neuronal loss.

The notion that BDNF limits the neuronal pathology of brain diseases such as AD, PD, depression, and schizophrenia has been extensively established (Zuccato and Cattaneo 2009). Yet, very little is known about a possible role of this neurotrophin in HAD, the most severe form of HAND. As presented above, AD (Kunugi et al. 2001; Phillips et al. 1991), PD (Howells et al. 2000; Nagatsu et al. 2000), depression, and other psychiatric disorders (Duman 2004; Post 2007; Toyooka et al. 2002) have in common a decrease in BDNF synthesis. Thus, while BDNF treatment could be a beneficial therapeutic approach in these diseases (Nagahara and Tuszynski 2011), at the same time, it is appealing to suggest that HIV may promote neuronal degeneration by lowering BDNF levels. This hypothesis has recently been tested in animal models of HAD as well as in HIV-positive subjects. It was found that HIV reduces BDNF both in rodents (Nosheny et al. 2004) and humans (Avdoshina et al. 2011). These data pose the question as to whether HIV promotes synaptic simplification by reducing BDNF or other neurotrophic factors whose physiological role is to support and maintain synapses (Fig. 1). This theory was first developed for sensory and sympathetic neurons in which the neurotrophins regulate the density of their innervation to a target cell. Consequently, the extent of neuronal loss is regulated by interactions with postsynaptic targets producing NGF (Johnson et al. 1980; Levi-Montalcini 1987). Nevertheless, such concept can also be applied to the CNS. Indeed, low levels of BDNF have been shown to impair the innervation of the cortex by serotonergic neurons (Lyons et al. 1999).

BDNF can avidly be transported in axons and released upon activation of neuronal activity (Conner et al. 1997; Marini et al. 1998). When such transport is interrupted, the release diminishes, and, consequently, neurons do degenerate as



**Fig. 1** Model illustrating how HIV induces neuronal degeneration. (a) Physiological environment. BDNF, released in an activity-dependent manner, promotes maintenance of neuronal architecture. BDNF can also signal back to the cell body to keep it from undergoing apoptosis. (b) Infected CNS. HIV infects the brain and sheds gp120. The viral protein binds to chemokine receptors and reduce BDNF trafficking and release. (c) Pathological brain. The lack of trophic support evokes short processes and elimination or pruning of processes that eventually culminate in apoptosis

BDNF profoundly affects neuronal homeostasis (Fig. 1) and the ability of neurons to counteract naturally occurring neuroinflammatory responses. This is applicable to different neuronal populations that express TrkB (Chao 2003), such as striatal (Zuccato et al. 2001), hippocampal (Erickson et al. 2011), and cortical (Mamounas et al. 1995; Xie et al. 2010) neurons. Conversely, BDNF inhibits neuronal *apoptosis* and cell loss in animal models of HAD both in vitro (Bachis et al. 2003) and in vivo (Nosheny et al. 2007). The neuroprotective property of BDNF raises the question of how this neurotrophin interacts with membrane receptors to avoid the destructive process of HIV. The next section reviews some experimental evidence that illustrates a unique molecular mechanism of BDNF action that might allow for the suppression of the main neurotoxic pathway of HIV. In order to fully comprehend this mechanism, a brief overview of how HIV infects cells is necessary.

#### 4 HIV in the Brain

#### 4.1 HIV and Chemokine Receptors

The most likely route for brain invasion by HIV is across the blood–brain barrier (BBB) (Price et al. 1988) although the choroid plexus and CSF pathway may also be

implicated (Petito et al. 1999). The primary CNS cell types that are vulnerable to HIV infection are the parenchymal microglia and the perivascular microglia/macrophages (Kramer-Hammerle et al. 2005). These monocyte lineage cells express the primary HIV receptor, CD4, a member of the immunoglobulin superfamily essential for activating helper cells, and chemokine co-receptors CCR5 or CXCR4 (Clapham and McKnight 2001). Chemokines represent a subset of cytokines with chemoattractant properties that are particularly important in HIV-related pathology. Indeed, both CXCR4 and CCR5 mediate entry of different strains of HIV into macrophages and T lymphocytes, respectively (Berger et al. 1999). Resident macrophages and microglia are the only indigenous CNS cell types to express both CD4 and CXCR4/CCR5, while neurons and astrocytes only express chemokine receptors (Albright et al. 1999). Consequently, macrophages and microglia are the primary target cells that are infected by a chemokine receptor-mediated mechanism (He et al. 1997). Infection of microglia by HIV has two potential effects: firstly, production of new virions and viral proteins and, secondly, induction of aberrant cytokine expression within the CNS (Anderson et al. 2002). Apart from microglia, astrocytes are also a possible target but are thought to be capable of only a restricted form of HIV infection in vivo (Brack-Werner 1999). The mechanism by which astrocyte infection occurs is unclear, as these cells do not express CD4, although, as stated above, they do express the chemokine receptors CCR5 and CXCR4. Recent reports have suggested a CD4-independent viral entry into astrocytes which is mediated either by mannose receptors (Liu et al. 2004) or through a gap junction mechanism (Eugenin et al. 2011). Regardless of whether astrocytes are infected or not, it is likely that these cells play a key role in the neuropathogenesis of HIV infection if their crucial functions in supporting neurons and maintaining BBB integrity are disrupted. These functions include the production of neurotrophic cytokines and buffering of substances such as glutamate which are neurotoxic at high concentrations.

There have also been occasional reports of neurons being infected with HIV, though neurons, like astrocytes, do not express CD4 but do express the chemokine receptors CCR5 and CXCR4 (Lavi et al. 1997; van der Meer et al. 2000). Given the lack of evidence for significant neuronal infection, it is thought that neuronal damage and death in AIDS is an indirect result of HIV infection of other cells in the brain. However, the mechanism(s) by which HIV infection of the CNS leads to neurocognitive disorders is not well understood and it is, therefore, the subject of intense research. The next chapters will briefly review some of the theories about HIV neurotoxicity that were obtained mostly from animal studies. The caveat is that simple animal models of AIDS/HAD needed to be carefully reevaluated in terms of relevance to the condition observed in humans.

#### 4.2 Viral Proteins

HIV may promote neuronal injury by distinct series of events. HIV infection causes *HIV encephalitis* (Everall et al. 2005) which is characterized by neuroinflammation, astrogliosis, and microgliosis and results in an overall production and release of

pro-apoptotic chemokines, such as interleukin-1 $\beta$  (IL-1 $\beta$ ) and tumor necrosis factor- $\alpha$  (TNF $\alpha$ ), and overexpression of CXCR4 (Mocchetti et al. 2008) and its ligand CXCL12 (Langford et al. 2002). These findings have led to the notion that activated microglia play a role in the pathogenesis of HIV encephalitis and suggested that neurodegeneration is triggered by host factors generated from the infection (Kaul et al. 2001). However, this theory does not take in account that activated (proliferating) microglia and released chemokines could be beneficial to recruit other cells in order to prevent any further damage to the infected brain (Cardona et al. 2006). An alternative mechanism of HIV-mediated neurodegeneration involves toxic proteins derived directly from the virus. The HIV genome codes for nine proteins that, apart from their roles in the viral life cycle, can damage neuronal cells and/or interfere with CNS function. One viral *protein* that can cause neuronal injury is the transactivator of transcription, or Tat, which has been shown to be released from HIV-infected cells (Ensoli et al. 1993). Tat can reduce neuronal survival by several indirect and direct mechanisms, among others the production of inflammatory cytokines such as tumor necrosis factor- $\alpha$ (Chen et al. 1997), impairment of mitochondrial function (Norman et al. 2007), activation of NMDA receptors (Haughey et al. 2001), and apoptosis (Kruman et al. 1998). These effects occur at concentrations lower than those needed to support viral replication. Additional viral proteins that evoke neuronal cell death include the accessory proteins Nef and Vpr (Li et al. 2005). These proteins are not released (Tornatore et al. 1994) but can cause neurotoxicity by a number of mechanisms including activation of potassium and calcium channels (Piller et al. 1996; Zegarra-Moran et al. 1999). Thus, a new theory of neurotoxicity has been suggested that calls for a direct interaction of these proteins with membrane-associated receptors and activation of a signaling pathway that promotes cell death.

One important viral protein that has been linked to the pathogenesis of HAD is the envelope protein gp120. This protein is necessary for infectivity as it allows the virus to change conformation after binding to chemokine co-receptors CXCR4 and CCR5 (Scarlatti et al. 1997). During viral entry gp120 can be shed from the virus and bind to CCR5 and CXCR4. This binding activates a cytotoxic pathway in both T cells and neurons (Grivel and Margolis 1999; Kaul et al. 2007). Indeed, activation of neuronal CXCR4 and CCR5 has been shown to promote neuronal loss/atrophy that is relevant to explain the neuropathology of HAD (Hesselgesser et al. 1998; Meucci et al. 1998; Zheng et al. 1999). Proof of this theory has also been obtained in transgenic mice expressing gp120 or in rats infused intracerebroventricularly with gp120. In fact, transgenic mice exhibit neuronal loss and *dendritic simplification* (Toggas et al. 1994), pathological features seen in postmortem brains of HAD subjects. Infusion of gp120 into rat brains causes apoptosis and neuronal loss (Acquas et al. 2004; Bagetta et al. 1995). Moreover, gp120 has also been shown to cause axonal degeneration (Melli et al. 2006) and dendritic injury (Everall et al. 2002; Iskander et al. 2004), two key pathological events that may account for the synaptodendritic atrophy observed in HAD (Masliah et al. 1997). These data are significant from the standpoint of experimental therapeutics because they allow the speculation that gp120 signal transduction could be a valid target for an adjunct therapy to prevent HIV neurotoxicity. Most importantly, they indicate that experimenters can recreate an animal model of HIV neurotoxicity without the need to deal with HIV infection.

# 5 Chemokine Receptors and Neurotrophins: From Experiments to Therapy

#### 5.1 Chemokine Receptors and Synaptic Injury

As described above, CXCR4 and CCR5 are crucial for gp120-induced neurotoxicity. Therefore, inhibitors of the expression/signaling of these receptors may have an important therapeutic property against HIV-mediated neurological impairment caused by CXCR4/CCR5 activation. These receptors are abundant in neurons and areas of the CNS that also express TrkB, such as the cortex, hippocampus, and striatum (Ahmed et al. 2008). CXCR4 and CCR5 are G-protein-coupled receptors that can be desensitized by tyrosine kinase intracellular signal crosstalk (Daub et al. 1996). Thus, TrkB activation may downregulate CXCR4 and CCR5. Indeed, experimental evidence has shown that BDNF and the other neurotrophins modulate the expression of CXCR4 and CCR5. This evidence includes the ability of BDNF to reduce CXCR4 in vitro (Bachis et al. 2003) and in vivo (Nosheny et al. 2007). Conversely, BDNF heterozygous animals exhibit increased levels of CXCR4 and CCR5 mRNA in the cortex, hippocampus, and striatum when compared to wild type (Ahmed et al. 2008). Moreover, in BDNF heterozygous mice, increased CXCR4 correlates with a more robust neurotoxic effect of gp120 (Nosheny et al. 2004), supporting the role of CXCR4 in gp120-mediated neuronal degeneration. Thus, from a pharmacological point of view, BDNF is particularly important as a neuroprotective compound against HIV or gp120 neurotoxicity that occurs through CXCR4 or CCR5 receptors.

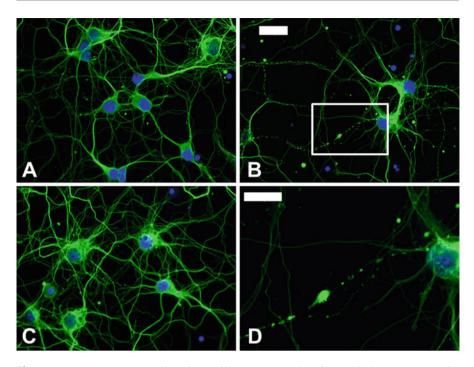
The neuroprotective effect of BDNF against gp120 relies on TrkB-mediated activation of the extracellular-signal-regulated kinase (ERK) pathway (Mocchetti and Bachis 2004). Conversely, BDNF does not prevent gp120 toxicity in neurons that do not express TrkB (Ahmed et al. 2008), suggesting that BDNF may not be suitable to reduce CXCR4 and HIV toxicity in all brain cells. However, from a therapeutic perspective, this is not an issue, as trophic factors can promote plasticity on overlapping as well as dissimilar neuronal populations/compartments. For example, cholinergic neurons of the basal forebrain express TrkB as well as TrkA. These neurons are more responsive to NGF than BDNF (Koliatsos et al. 1994). Thus, NGF could be used to modulate the plasticity of cholinergic neurons and control their function in learning and memory in HAD. Moreover, basic fibroblast growth factor, another neurotrophic factor that binds to tyrosine kinase receptors and has been shown to downregulate CXCR4 (Sanders et al. 2000), could also be used to rescue Trk-negative neurons and therefore overcome the toxic effects of HIV in neurons that do not respond to the neurotrophins. Based on all the experimental data collected so far, it is clear that specific neurotrophins may be necessary to stimulate growth in a defined population of CNS neurons. The functional significance of the ability of the neurotrophins to modulate CXCR4 remains to be established as well as whether neurotrophins act on more than one chemokine receptor.

The survival of neurons exposed to HIV is greatly enhanced by CXCR4 or CCR5 receptor antagonists (Bachis et al. 2009). Moreover, as discussed before, several groups have demonstrated that gp120 binding to chemokine receptors evokes caspase-3-mediated apoptosis. This is relevant for HAD because cortical neurons of HIV subjects are caspase-3 positive (Garden et al. 2002). Thus, the ability of BDNF to prevent gp120 activation of caspase-3 (Bachis et al. 2003) suggests that this neurotrophic factor will also be neuroprotective against HIV. However, postmortem studies of HAD have also indicated that neuronal apoptosis does not correlate with dementia and that loss of dendritic spines and synaptic densities may be the primary cause leading to apoptosis (Adle-Biassette et al. 1999). Consistent with this hypothesis, gp120 reduces dendritic arborization and dendritic volume in human neurons in vitro (Iskander et al. 2004). However, such pathological changes can also be reproduced in rodent neurons in vitro by HIV. In fact, a dramatic alteration of axonal processes, characterized by fragmentation and varicosities, can be seen as quickly as 6–8 h after exposure to HIV (Fig. 2). Most importantly, the axonal disruption is preventable by BDNF. Thus, BDNF can also help neurons to survive by inhibiting the disruption of axons caused by HIV.

#### 5.2 Delivery of BDNF

Of particular theoretical importance is the use of BDNF in HIV-positive patients as an adjunct therapy combined with cART. As discussed above, the physiological role of BDNF in promoting the survival and plasticity of adult neurons adds credence to the experimental data supporting such therapy. Moreover, in vitro models of HIV neuropathogenesis have indicated that HIV infection of the brain triggers the release of glutamate (Bezzi et al. 2001). Glutamate, through excitotoxicity, may help propagate the initial "injury" caused by inflammation to adjacent neurons. BDNF is a strong inhibitor of *excitotoxicity* whether it is induced by neurotoxins (Marini et al. 1998; Martinez-Serrano and Bjorklund 1996) or ischemia (Han and Holtzman 2000). Thus, BDNF may also be a potential therapeutic agent in preventing HIV-mediated excitotoxicity, especially considering the side effects of glutamate receptor antagonists.

A crucial question to be considered is what constitutes a safe way of administering BDNF into the CNS. A major concern is the degree of penetration of BDNF into the CNS. BDNF is a charged protein that does not efficiently cross the BBB. When delivered by direct infusion into the ventricles, its parenchymal diffusion is limited; thus, the delivery must be constant. More recently, viral vectors have been designed to deliver the gene encoding for BDNF into the CNS (Klein et al. 1998). These vectors, such as *adeno-associated viruses* (AAVs), use viral backbones to transduce host cells in a brain area of interest. AAVs are safe and capable of sustaining long-term BDNF expression in neurons (Klein et al. 1998). Using an AAV-expressing BDNF, it was shown that BDNF prevents gp120-mediated



**Fig. 2** *BDNF prevents HIV-mediated axonal damage.* Examples of rat cortical neurons grown for 7 days in vitro and then exposed for 6 h to (**A**) control medium (0.1 % BSA); (**B**) medium containing HIV; (**C**) medium containing BDNF and HIV; and (**D**) higher magnification of the *square* shown in (**B**) to show typical axonal varicosities. *Blue* DAPI to visualize cells, *green*  $\beta$ III-tubulin to visualize neuronal processes. *Bar:* A, B, and C = 20 µm and D = 10 µm. Immunofluorescence was analyzed using a FV300 laser confocal scanning system attached to an Olympus IX70 upright microscope. Please note the lack of axonal swelling and varicosities, the hallmarks of diffuse axonal injury (Tang-Schomer et al. 2010), in BDNF-treated cultures

neuronal apoptosis in vivo (Nosheny et al. 2007), raising the possibility of using BDNF in clinical trials in the near future. However, caution must be exerted as clinical effects could be accompanied by undesired effects. For instance, BDNF also promotes sprouting of dendrites of dopaminergic neurons in gp120-treated rats (Nosheny et al. 2007). Although, in general, this is a trophic effect of the neurotrophins on neuronal processes, it may be a negative side effect of overdosing. Indeed, according to the classical view of trophic molecules, a trophic factor must be delivered in small amounts and only to specific populations of neurons to avoid such side effects.

#### 5.3 BDNF and Immune Responses

There are other effects of BDNF that can be beneficial for HIV-positive subjects. For instance, in the immune system, BDNF decreases apoptosis of T cells

(De Santi et al. 2009; Maroder et al. 1996). These cells are depleted in AIDS. Thus, one may envision the use of BDNF in conjunction with cART to maintain the appropriate number of immune cells and delay AIDS. This effect also has a broader implication for the CNS function. In fact, as progressive neurological deficits occur after the onset of severe immunodeficiency (McArthur et al. 2010), keeping healthy immune cells will reduce "inflammation" of the immune system that can amplify nervous system damage via inflammatory cytokines. These may enter the CNS either from the periphery or produced locally by *microglia*. For example, in a state of chronic inflammation induced in a nonhuman primate by infection with simian immunodeficiency virus, the brain was positive for peripheral circulating monocytes trafficking from bone marrow. This event correlates with the severity of encephalitis (Burdo et al. 2010). Abnormal activation of cytokine and chemokine receptors in the context of HIV infection results in dendritic

beading and loss of dendritic spines. These changes are accompanied by failure of long-term potentiation (LTP), which might underlie impaired learning and memory. Given the well-known property of BDNF in promoting dendritic branching and spine morphology (Horch and Katz 2002; Tanaka et al. 2008) which are key for the BDNF-mediated LTP (Figurov et al. 1996; Gartner and Staiger 2002), BDNF could be used to prevent atrophy of *dendrite branching* and memory loss. Of course such treatment will require an early diagnosis of cognitive impairments.

## 6 Conclusion

The dividing line between viral- and host-mediated neurotoxicities in HIV infection is neither precise nor rigid. An effective therapy aimed at treating HIV-induced CNS diseases will require both cART and adjunctive therapies that include neuroprotective and neuro-regenerative agents. Neurotrophic factors are large hydrophilic molecules that do not cross the BBB. Therefore, these molecules need to be delivered within the CNS, which lowers the enthusiasm for a chronic instrumentation in a nonclinical setting. In addition, when infused into the CNS, BDNF has limited diffusion. The delivery problems may not be severe for AIDS patients with neurodegeneration of peripheral neurons (i.e., peripheral neuropathy) as these neurons are located outside the CNS and are capable of retrogradely transporting BDNF to the soma. However, the efficiency of terminal uptake and axonal transport of trophic molecules is uncertain as there is evidence of impaired axonal transport in AIDS. One strategy, still under experimentation in humans, is the delivery of BDNF via viral vectors. A major concern about a clinical approach to increase BDNF in the CNS by this method is that an abnormal overproduction of BDNF may cause side effects that may be intolerable over a long period of time. Such effects may be seen locally (fiber spouting around the site of BDNF production) or systemically (weight loss). Thus, additional strategies must be applied to circumvent these problems. Ongoing research has demonstrated more safe and reliable methods to increase endogenous BDNF and consequently influence neuronal survival. This could replace the need for delivering BDNF by invasive approaches. For instance, physical exercise has been reported to increase BDNF levels in the hippocampus (Cotman and Berchtold 2002) which, in turn, promotes hippocampus-based learning, synaptic plasticity, and neurogenesis (Vaynman et al. 2004). Antidepressant drugs, which are safely used in humans, also increase BDNF synthesis (Hashimoto et al. 2002; Nibuya et al. 1996). Thus, there are alternatives to minimize the use of chronic instrumentation.

Besides its traditional role as a molecule promoting survival and regeneration, BDNF is seen as a potent mediator of synaptic plasticity in the adult CNS. These effects could play a role in limiting AIDS-mediated neuronal degeneration and stimulating neuronal function. Although still tested only in animal models of HAD, the powerful properties of BDNF provide incentive for further research and raise hope for the therapeutic possibilities in the near future.

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# **Neurotrophic Therapy for ALS/MND**

# Mary-Louise Rogers

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#### Abstract

Neurotrophic factors have a large range of activities in the nervous system that consist of functions in development, plasticity, neurogenesis, disease, and injury. In the context of amyotrophic lateral sclerosis (ALS), it has long been hypothesized that *lack* of neurotrophic growth factors *is one of the neurotoxic* contributors to the

e-mail: mary-louise.rogers@flinders.edu.au

M.-L. Rogers

Department of Human Physiology, Centre for Neuroscience, School of Medicine, Flinders University, Adelaide, SA, Australia

disease that results in death of motor neurons. This has led to a considerable number of clinical trials undertaken involving neurotrophic therapy for ALS, although none have shown benefit. This chapter will review the cause and pathology of ALS and how neurotrophic factors relate to neurotoxicity in this disease. The treatments targeted at neurotoxicity and results of trials will be discussed, in particular neurotrophic factors. This will include glial cell-derived neurotrophic factor (GDNF), brain-derived growth factor (BDNF), neurotrophin-3 (NT-3), ciliary neurotrophic factor (CNTF), insulin-like growth factor (IGF), vascular endothelial cell growth factor (VEGF), fibroblast growth factor (FGF), and hepatocyte growth factor (HGF). Also highlighted is the potential for reexamining neurotrophic factors as treatments for ALS, including new delivery methods.

Keywords	
ALS • BDNF •	$CNTF \bullet FGF \bullet GDNF \bullet IGF \bullet NGF \bullet NT-3 \bullet NT-4 \bullet NT-4 \bullet VEGF$

List of Abbrevi	ations
ALS	Amyotrophic lateral sclerosis
BBB	Blood–brain barrier
BDNF	Brain-derived growth factor
CNTF	Ciliary neurotrophic factor
CNS	Central nervous system
FGF	Fibroblast growth factor
GDNF	Glial cell-derived neurotrophic factor
HGF	Hepatocyte growth factor
IGF	Insulin-like growth factor
NGF	Nerve growth factor
NMJ	Neuromuscular junction
NT-3	Neurotrophin-3
NT-4	Neurotrophin-4
NT-5	Neurotrophin-5
SOD1	Superoxide dismutase 1
SOD1G93A	Mutated SOD1 glycine to alanine position 93
TDP-43	Transactivation response DNA-binding protein of 43 kDa
VEGF	Vascular endothelial cell growth factor

# 1 Introduction: What Are the Neurotrophic Factors and How Do They Relate to Neurotoxicity in ALS?

The story of neurotrophic growth factors in the nervous system began when Rita Levi-Montalcini, Viktor Hamburger, and Stanley Cohen discovered the actions of nerve growth factor (NGF) in the 1950s (Levi-Montalcini and Hamburger 1951; Levi-Montalcini and Cohen 1960). In 1986, both Levi-Montalcini and Cohen

received the Nobel Prize for the "discovery of growth factors" and at that time were already acknowledged for opening up a large area of research and potential therapeutic approaches for these agents within and outside the nervous system. The list of proteins regarded as neurotrophic growth factors has grown, and a lot has been learned about common and specialized activities of these factors. Some of them, the four-member neurotrophin family that includes NGF, were first described as acting on neuronal cells, whereas other more recently described neurotrophic factors were initially discovered in a nonneuronal context.

Neurotrophic factors have an array of activities in the nervous system that consist of functions in development, plasticity, neurogenesis, disease, and injury. In the context of amyotrophic lateral sclerosis (ALS), it has long been hypothesized that *lack* of neurotrophic growth factors *is one of the neurotoxic* contributors to the disease that results in death of motor neurons. This has led to a considerable number of clinical trials undertaken involving neurotrophic therapy for ALS, although none have shown benefit. In retrospect this may be due to inadequate dosing and/or route of application of the growth factor in question. This chapter will review the general contributors to neurotoxicity in ALS including withdrawal of neurotrophic support and then the therapies trialled which are aimed at correcting neurotoxicity including neurotrophic therapy, highlighting the potential for reexamining neurotrophic factors as treatments.

The first neurotrophic factor to be discovered was NGF (Levi-Montalcini and Hamburger 1951). Subsequent systematic search of mammalian tissues revealed that male mouse salivary glands contained substantial amounts of NGF (Cohen 1960; An odd finding that has never been clearly understood!). Thirty years after the discovery of NGF, another neurotrophin that demonstrated neuronal growthpromoting properties was purified from pig brain, and later called brain-derived neurotrophic factor (BDNF; Barde et al. 1982). A third member of the family called neurotrophin 3 (NT-3) was identified in 1990 (Hohn et al. 1990) and neurotrophin 4 (NT-4) in 1991 (Hallböök et al. 1991), with neurotrophin 5 (NT-5) later identified as identical to NT-4 (Ip et al. 1993). After the discovery of NGF, a number of other, unrelated neurotrophic factors in the nervous system were discovered that have a similar scope of activities, including ciliary neurotrophic factor (CNTF; Barbin et al. 1984; Lin et al. 1989) and the glial cell line-derived neurotrophic factor (GDNF; Lin et al. 1993). Members of the insulin-like growth factor (IGF; Shemer et al. 1987; Rotwein et al. 1988) fibroblast growth factor (FGF; (Emoto et al. 1989)) and vascular endothelial growth factor (VEGF; Alvarez et al. 1992) families are known to have roles in the nervous system additional to those outside the nervous system. In addition, hepatocyte growth factor (HGF), which was found in the liver, has further actions in the nervous system (Jung et al. 1994).

## 2 Searches for the Cause and Pathophysiology of ALS

In 1869, the French neurologist Jean-Martin Charcot described a fatal and progressive paralysis that he termed amyotrophic lateral sclerosis (ALS) (Rowland 2001). This devastating neurodegenerative disease, also known as Lou Gehrig's disease in the USA or motor neuron disease in the UK and Australia, has a worldwide incidence of 2 per 100,000 and prevalence of 7 per 100,000 (Ferraiuolo et al. 2011). ALS is characterized by progressive, debilitating paralysis. Patients experience muscle weakness and atrophy, leading to inevitable paralysis, due to the selective and progressive loss of motor neurons of the cerebral cortex, brain stem, and spinal cord that is characteristic of the disease. Most often, patients succumb to the disease because of denervation of muscles involved in respiration (Vande Velde et al. 2011). ALS/MND occurs mainly in adults (45–60 years of age), and only 50 % of patients survive 2–3 years after diagnosis. There is only one therapeutic option for patients, riluzole, which offers a modest 3- to 6-month extension in survival. However, in the last 3 years, real progress has been made in the identification of genetic causes of the disease, such as expanded GGGGCC hexanucleotide repeats in a noncoding region of C9ORF72 (Renton et al. 2011; Zinman and Cudkowicz 2011; Ludolph et al. 2012).

Around 5–10 % of ALS patients have inherited a mutation that can cause familial ALS (Andersen and Al-Chalabi 2011), with the remaining 90 % developing the disease sporadically through environmental and lifestyle risk factors (Sutedja et al. 2009). The highest proportion of familial ALS patients identified is that with expanded GGGGCC hexanucleotide repeats in the noncoding region of C9ORF72 (Renton et al. 2011), followed by patients that carry Cu/Zn superoxide dismutase 1 (SOD1), mutations (Ludolph et al. 2012). Other rarer mutations leading to ALS occur in genes such as alsin, senataxin, dynactin, transactivation response DNA-binding protein of 43 kD (TDP-43), fused in sarcoma/translated in liposarcoma (FUS), optineurin, and ubiquilin 2 (Deng et al. 2011; Andersen and Al-Chalabi 2011; Ludolph et al. 2012). Over 160 SOD1 mutations have been identified (Gurney et al. 1994; Al-Chalabi et al. 2012) in humans.

Transgenic rodents carrying SOD1 mutations have been essential to understanding the mechanisms of ALS in humans (Peviani et al. 2010; Kanning et al. 2010). All mimic human familial and sporadic ALS, with a dominant gain of toxic function resulting in motor symptoms and death, due to the neurotoxicity of the mutant SOD1 protein (Reaume et al. 1996; Turner and Talbot 2008; Peviani et al. 2010; Kanning et al. 2010). Transgenic mice that carry 23 copies of the human mutant SOD1G93A transgene are used for preclinical testing of ALS treatments (Scott et al. 2008; Ludolph et al. 2010). Mice that overexpress normal or human mutant TDP-43 have been produced (Cohen et al. 2011), but they have not yet lived up to their promise as a model for ALS. Several groups have generated TDP-43 knockout mice and in each instance all die during embryonic life, indicating the importance of TDP-43 in multiple biological processes (Wu et al. 2010; Kraemer et al. 2010). Issues with mice expressing mutant TDP-43 have been identified such as inconsistent pathology and disease progression across TDP-43 mouse models and most recently the TDP-43 A315T mutant mice dying from neuropathy-linked bowel obstruction (Igaz et al. 2011; Guo et al. 2012). Conversely, evidence supports the continued relevance of SOD1G93A animal models to sporadic disease. Misfolding of SOD1 into a disease-specific and toxic conformation has been found in both sporadic and familial ALS (Bosco et al. 2010). Moreover, SOD1G93A mice replicate features of human MND including progressive limb paresis, respiratory failure, and reduced life span (Gurney et al. 1994; Turner and Talbot 2008). Behavioral symptoms appear around 90–110 days, but pathological changes occur much earlier (Peviani et al. 2010; Kanning et al. 2010).

Although we are yet to understand how the disease initiates, there is a large body of data on mechanisms that cause the pathology in motor neurons such as oxidative stress, protein aggregation, excitotoxicity, impairment of axonal transport, mitochondrial dysfunction, altered DNA/RNA processing, endoplasmic reticulum stress, and apoptosis (reviewed in Ferraiuolo et al. 2011). A common feature is ubiquitin-positive inclusions, most of which contain TDP-43 (Geser et al. 2011) that does not appear to be related to TDP-43 mutations. The pathophysiology of the disease includes a reduced secretion of neurotrophic factors, protein aggregations, malfunctioning of the mitochondria, rupture in the axonal passage, destruction in the calcium metabolism, changes in the skeletal proteins, high levels of glutamate, and oxidative damage (Ferraiuolo et al. 2011). Preventing and slowing down motor neuron degeneration and death in ALS are critical goals of future therapies, as are means of enhancing axonal regeneration.

Axonal transport has long been implicated in ALS. Axonal transport uses microtubule-dependent kinesin and cytoplasmic dynein molecular motors, which mediate transport towards the neuromuscular junction (anterograde transport) and towards the cell body (retrograde transport), respectively. Given the extremely long axons of motor neurons and their dependence on axonal transport for the delivery of ATP-producing mitochondria and other essential constituents, including neurotrophic factors, it comes as no surprise that transport deficiencies cause debilitating effects. Tumor necrosis factor elevated in SOD1G93A mice disrupts kinesin function via a mechanism involving p38 mitogen-activated protein kinase (p38 MAPK; De Vos et al. 2000), activation of which has been demonstrated in models of ALS (Holasek et al. 2005; Dewil et al. 2007). Mutant SOD1 also inhibits anterograde fast axonal transport of mitochondria and enhances the retrograde movement resulting in a depletion of axonal mitochondria content (De Vos et al. 2007). A study suggested that alterations in retrograde signalling contribute to neurodegeneration in ALS, by shifting from the transport of survival-promoting to death-promoting signalling cargo molecules (Perlson et al. 2009), suggesting less neurotrophic factors can be transported. Point mutation of the p150 subunit of dynactin has been detected in both sporadic and familial ALS patients (Munch et al. 2004). Mutations in dynactin lead to a reduction in retrograde transport in SOD1G93A mice (Puls et al. 2003).

Peripherin and neurofilament are intermediate filament (IF) proteins that can be abnormally found in inclusion bodies of the perikaryon and axon of motor neurons in ALS. Rare mutations in peripherin have been identified in a small number of people with ALS (Gros-Louis et al. 2004; Corrado et al. 2011). In addition to being found in inclusion bodies, neurofilaments are transported by axons and disruption of neurofilament transport occurs in ALS. Neurofilaments comprise three major proteins, neurofilament light, middle, and heavy chains (NFL, NFM, and NFH), that are synthesized within cell bodies and then usually transported into and through axons by "axonal transport." Decreased amounts of normal NFH in axons of SOD1G93A mice are correlated to shrinkage of axons in the ventral root and precede symptoms in SOD1G93A mice (Zhang et al. 1997). Abnormal phosphorvlation of the side-arm domain of NFH by the stress-activated kinase p38alpha has been reported in sporadic and familial ALS and SOD1G93A mice (Ackerley et al. 2000, 2004). This work indicated the speed of neurofilament transport through axons is altered (Ackerley et al. 2004) with a slowing of transport proportional to the increase in phosphorylation (Ackerley et al. 2000). Excitotoxic damage by glutamate is thought to contribute to motor neuron injury in ALS. In cultured neurons, glutamate reduces axonal transport of neurofilaments via activation of protein kinases that phosphorylate NFM such as p38 (Ackerley et al. 2000), and riluzole protects against glutamate-induced slowing of neurofilament axonal transport (Stevenson et al. 2009). An association of neurofilaments with ALS is also represented by rare mutations in the neurofilament gene in ALS patients (Al-Chalabi et al. 1999). There is a significant increase in NFM in CSF and serum of humans with ALS and SOD1G93A mice (Brettschneider et al. 2006; Ganesalingam et al. 2011), especially in the late stage of disease (Lu et al. 2012). Thus, axonal transport defects are an important feature of ALS that may precede motor neuron degeneration and clinical symptoms.

At the tissue level, it is clear that cells beyond motor neurons are involved in ALS pathophysiology (Ilieva et al. 2009). Cells such as glia and immune cells are proposed to be involved in non-cell-autonomous toxicity. For example, astrocytes are reactive and apparently toxic in ALS (Ince et al. 2011). Also, reactive microglia and T-cell infiltration have been noted as players in non-cell-autonomous toxicity (Ilieva et al. 2009). Most of this data has come from studies in which human mtSOD1 is expressed in rodents. There is evidence that the onset of ALS in mutant SOD1 mice results from mutant SOD1 action in the motor neurons, whereas progression of the disease is a result of the action of SOD1 mutant protein in glia (Ilieva et al. 2009). Selective reduction of mutant SOD1 from astrocytes or microglia using cre-lox system slowed disease progression in SOD1G37R mice (Yamanaka et al. 2008; Boillee et al. 2006). A complimentary approach replacing microglia/macrophage via bone marrow transplantation demonstrated that wild-type microglia/macrophage slow disease progression in SOD1G93A mice although inflammation may be a consequence, not an initiator of disease (Kano et al. 2012).

Advances in identification of upstream mechanisms that might contribute to the death of motor neurons in ALS and compounds directed at these potential targets are being developed. Ongoing research is addressing the question of where the motor neuron disease process is initiated, candidates being the neuro-muscular junction (NMJ), the axon, mitochondria, proteasome, or the cell nucleus (Dupuis and Loeffler 2009; Bilsland et al. 2010; Shi et al. 2010; Dupuis et al. 2004; Baron et al. 2007; Daoud and Rouleau 2011; Keller et al. 2012). Progress in understanding this etiology has led to therapies being trialled to target the various candidates that are causative in the disease. In addition, although ALS is a motor neuron disease, interventions directed towards improving muscle function such as muscle troponin activators, myostatin inhibitors, and reticulon 4 (Nogo-A) inhibitors are also proposed as therapies (Zinman and Cudkowicz 2011).

# 3 Treatments for ALS Targeted at Neurotoxicity

# 3.1 Treatments Targeted at Downregulation of SOD1 Mutations

For the approximately 20 % of familial patients with SOD1 mutations leading to familial ALS, a phase 1 human trial of safety, tolerability, and pharmacokinetics of antisense SOD1 oligonucleotides administered intrathecally to patients with SOD1 familial ALS is underway (ISIS 333611/clinicaltrials.gov/NCT01041222). The trial enrolled 21 patients with SOD1 mutations who were randomized into either three ascending dose cohorts or a placebo group. Some patients elected to enroll a second time. The phase 1 trial was completed in Jan 2012 with no significant adverse effects (Robinson 2012). However, some concerns were raised that the antisense oligonucleotide used in the trial targeted the mRNA of the non-mutated portion of SOD1 and that there may be a risk of inflammation in long-term, high-dose treatments. Isis the pharmaceutical company involved in this trial has recommended further animal trials with a redesigned antisense SOD1 oligonucleotide (Robinson 2012). The development of vaccine or passive infusion delivery of immunoglobulin to remove misfolded protein in ALS is a novel therapeutic strategy under investigation. Bosco et al. (2010) found misfolding of SOD1 into a disease-specific and toxic conformation underlies both sporadic and familial MND. Subsequently, vaccination with SOD1 mutant protein or antigenic peptide directed to monomer/misfolded SOD in the ALS SOD1 transgenic mouse model delayed disease onset and significantly extended survival (Takeuchi et al. 2010; Liu et al. 2012).

## 3.2 Treatments Targeted at Excitotoxicity

Glutamate-mediated excitotoxic effects have long been postulated to have an important role in motor neuron degeneration, with riluzole an example of a glutamate inhibitor used to treat ALS patients (Bensimon et al. 1994; Lacomblez et al. 1996). In an ALS SOD1 mouse model of ALS, lithium showed reduced aggregates of alpha-synuclein, ubiquitin, and SOD1 in motor neurons, and in addition to its known effects on protein clearance, lithium was found to increase the number of mitochondria and suppressed reactive astrogliosis (Fornai et al. 2008). However, lithium by itself failed in human clinical trials (Verstraete et al. 2012), and more recently in combination with riluzole, although well tolerated had no benefit (Aggarwal et al. 2010). Arimoclomol is an oral compound that amplifies the heat-shock response leading to induction of endogenous cellular cytoprotective mechanisms (Kieran et al. 2004) that might prevent motor neuron degeneration under conditions of stress. Arimoclomol delayed disease progression and extended survival by 22 % in the SOD1G93A transgenic ALS mice (Kieran et al. 2004; Kalmar et al. 2008; Lu et al. 2012). Efficacy studies in patients with familial ALS are underway with promising early results indicating this drug is safe in human beings, and it crosses the blood-brain barrier (BBB) following oral administration (Cudkowicz et al. 2008).

# 3.3 Treatments Targeted at Mitochondria

Results from ongoing trials using dexpramipexole, a drug that improves mitochondrial health, showed initial promise in ALS trials. Dexpramipexole is an optical enantiomer of pramipexole, a Parkinson's disease drug that maintains prolonged survival of ALS SOD1 transgenic mice (Danzeisen et al. 2006). A large, Phase 3 international study of treatment of ALS patients with dexpramipexole (EMPOWER) has been completed with disappointing results. There was initial promise with the first reports from a carefully controlled Phase 2 study indicating subjects show a slower rate of decline in measures of motor function and a lower level of mortality over the course of the study (Bozik et al. 2010; Cudkowicz et al. 2011: Rudnicki et al. 2013). However, in January 2013, the Phase 3 trial was suspended due to lack of effect (alsa.org/news/archive/biogen-discontinues.html). Some compounds proposed to improve mitochondrial function, such as minocycline and creatine, have had beneficial effects in the SOD1 transgenic mouse model (Zhu et al. 2002) but have yet to prove beneficial in human trials (Klopstock et al. 2011; Gordon et al. 2007). A new trial of creatine in combination with tamoxifen is underway (www.clinicaltrials.gov: NCT01257581b). Other agents targeting mitochondrial function have been identified. Olesoxime (previously TRO19622) is a mitochondrial pore modulator that improves SOD1G93A mice survival and attenuates activation of surrounding glial cells. A phase 2/3 study of olesoxime is underway in Europe (NCT00868166).

# 4 Neurotrophic Factor Withdrawals and Neurotrophic Factors as Treatments

# 4.1 Rationale

ALS is a complex disease and cannot be explained by the occurrence of a single event or the disturbance of a single gene or protein. The disease is likely to be the result of a multistep process, ultimately leading to motor neuron degeneration and clinical symptoms. Currently there are two hypotheses to explain the etiology of ALS. The first is the "dying-forward" hypothesis which proposes that ALS is mainly a disorder of cortico-motor neurons, which connect monosynaptically with anterior horn cells, mediating anterograde degeneration of anterior horn cells via glutamate excitotoxicity (Kiernan et al. 2011). The second is the "dying-back" hypothesis where pathological changes in motor axons and nerve terminals appear to precede motor neuron degeneration and clinical symptoms. Moreover, this second hypothesis raises the possibility that the disease process may start distally at the nerve terminal or at the neuromuscular junction (NMJ) and progress towards the cell body (Dadon-Nachum et al. 2011). The main therapeutic strategy for ALS directed to the dying-back hypothesis is the use of neurotrophic factors.

No genetic defects have been identified to date that interfere with any of the growth factor genes in ALS. The lack of trophic support appears to be a downstream

consequence of axonal transport of neurotrophic factors in ALS rather than expression. A deficiency of growth factor support could provoke motor neuron death in patients with ALS. In early stages of the disease, nerve terminals and neuromuscular junctions are partially degraded, while cell bodies in the spinal cord are mostly intact. This is clearly shown by Fischer and colleagues (Fischer et al. 2004) who demonstrated, in autopsy material from an ALS patient who died from an unrelated cause only 6 months after symptom onset, that denervation and reinnervation were pronounced, while the spinal motor neurons were not yet affected. The concept is further supported by recent work showing progressive motor axonal changes play a significant role in the course of disease including clinical impairment (Cheah et al. 2012). Studies in animal models of spinal muscular atrophy (SMA), a pure lower motor neuron disease, have shown that the first event to be observed is withdrawal of the axon terminals from the adjacent muscle fibers (Simon et al. 2010). In SOD1G93A mice axonal degeneration also precedes onset of symptoms as well as spinal motor neuron loss. Disease progression in these animals follows the dying-back principle characterized by an ascending neuronal degeneration from the terminals of the motor axons to the motor neuron cell body (Gould et al. 2006). Retrograde transport of neurotrophic factors such as BDNF and GDNF is impaired, whereas death-promoting signalling molecules are enhanced in axons of motor neurons of presymptomatic SOD1G93A mice (Perlson et al. 2009). Both retrograde transport and anterograde axonal transport are impaired in human ALS and in mouse models. This is supportive of the dying-back hypothesis that leads to loss of muscle function and the inability of neurotrophic factors, such as glial cellderived neurotrophic factor (GDNF), to be transported to the motor neuron cell body.

# 4.2 Glial Cell-Derived Neurotrophic Factor

GDNF can be efficiently transported by motor neurons both retrogradely (Leitner et al. 1999) and anterogradely (Rind and von Bartheld 2002). In a study where stem cell-derived motor neurons were transplanted into paralyzed rat spinal cord and GDNF-secreting cells implanted into the sciatic nerve, axonal outgrowth and attachment to the muscle could be achieved. This thus supports the notion that combining protection of motor neuron cell body and support of nerve terminals in the muscle by GDNF may be beneficial in ALS (Deshpande et al. 2006). A study examining the overexpression of GDNF either centrally (astrocytes) or peripherally (muscle) showed that overexpression of GDNF in skeletal muscle, but not in the central nervous system (CNS), improved survival in the SOD1G93A model of ALS (Li et al. 2007). When stem cell-derived GDNF was expressed in astrocytes at disease onset, there was no improvement in survival in SOD1G93A mice although there was some improvement in health of motor neurons (Park et al. 2009). The outcomes of studies in SOD1G93A mice argued for a benefit of GDNF treatment. GDNF-treated motor neurons of SOD1 animals are clearly protected (Mohajeri et al. 1999; Acsadi et al. 2002; Suzuki et al. 2007). Some studies report a benefit for disease progression and/or survival (Acsadi et al. 2002; Li et al. 2007). Others

found GDNF has no effect on disease progression and integrity of neuromuscular junctions (Suzuki et al. 2007; Park et al. 2009), although Park et al. (2009) were the only ones to initiate treatment at the onset of disease.

Although GDNF has been used in clinical trial for Parkinson's disease, the factor has not yet been tested clinically for the treatment of ALS patients. Clinical observations demonstrated that the level of GDNF is increased in the CSF of ALS patients (Grundstrom et al. 2000) and that its receptor is expressed in the spinal cord, even at late stages of the disease (Mitsuma et al. 1999). Due to its in vivo neuroprotective effects on motor neurons, GDNF remains a potential candidate for a clinical trial in ALS, despite controversial efficacy data on survival. However, as there is evidence that GDNF does not cross the blood-brain barrier (Kastin et al. 2003), a future challenge remains in developing a protocol to resolve this issue, such as using an intrathecal delivery device for GDNF. Intraventricularly, GDNF has been trialled as a treatment for Parkinson's disease, but a clinical trial was halted because of adverse effects. Subsequent trials involving continuous intraputamenal infusion had no adverse effects but are inconclusive (Yasuda and Mochizuki 2010). Alternate routes for GDNF delivery include coupling to proteins that do cross the BBB. For example, intravenous, nonviral GDNF gene therapy in a rat model of Parkinson's disease was highly effective because the nonviral agent contained antibodies to the transferrin receptor that allowed entry across the BB (Zhang and Pardridge 2009). In another example, delivery of plasmid GDNF in neonatal rats was achieved using a nonviral complex containing antibodies to p75NTR receptor that enabled retrograde entry to motor neuron cell bodies from terminals in muscles (Barati et al 2006). GDNF gene therapy via direct delivery into the brain using a viral vector (AAV2-GDNF) is underway (clinicaltrials.gov).

# 4.3 Brain-Derived Neurotrophic Factor (BDNF)

BDNF was originally isolated as a trophic factor for neurons from brain extract (Barde et al. 1982). Expression of BDNF (Henderson et al. 1993) and the highaffinity BDNF receptor tropomyosin-related kinase B receptor (TrkB; Yan et al. 1993) is observed in skeletal muscle and motor neurons during motor neuronprogrammed cell death. BDNF/TrkB kinase signalling is required for maintenance of the cholinergic phenotype of adult motor neurons and prevents embryonic and postnatal motor neuron cell death in a variety of experimental paradigms. For example, BDNF rescues motor neurons from embryonic-programmed cell death and neonatal axotomy-induced death (Oppenheim et al. 1992; Yan et al. 1992; Koliatsos et al. 1993). Research has shown BDNF and NT-3 slow motor neuron degeneration and increase lifespan in two models of inherited motor neuron disease, the wobbler and progressive motor neuropathy (pmn) mice (Ikeda et al. 1995; Haase et al. 1997). BDNF can also protect neurons from in vivo excitotoxicity (Bemelmans et al. 2006), a situation relevant to ALS.

In several clinical trials BDNF has failed to show any beneficial effect for ALS. A trial where 1,136 patients were given  $25-100 \text{ }\mu\text{g/kg/day}$  subcutaneously

for 9 months was reported in 1999 (The BDNF Study Group 1999). In this trial, the tolerability of BDNF was shown to be acceptable, despite some adverse effects such as diarrhea. While the results failed to show an effect on survival after treatment with BDNF, it appeared that a subgroup of patients responded positively to the treatment and demonstrated a higher survival rate. This subgroup consisted of patients having the lowest vital capacity and receiving the highest dose of BDNF, suggesting that BDNF could be beneficial for patients in advanced stages of the disease. The investigators suggested use of higher doses or intrathecal delivery to deliver BDNF to the diseased spinal cord. 1 year later, a phase I trial was published, showing the feasibility of intrathecal delivery of BDNF. Indeed, delivering BDNF with this method showed that BDNF leads to a detectable level of BDNF in the CSF, especially at the lumbar level (Ochs et al. 2000). The method being feasible, two clinical trials were conducted in 2003 and 2005. In 2003, 11 ALS patients were enrolled in a trial to receive an intrathecal delivery of BDNF. Another trial with intrathecal delivery was conducted in 2005 with 13 patients (Kalra et al. 2003; Beck et al. 2005). Both trials were too small to detect any meaningful efficacy signal.

What have we learned? Failure of BDNF therapy has been attributed to factors such as short half-life, inadequate dosage, and limited bioavailability. The first issue is the short half-life; BDNF has a half-life of 0.92 min in the circulation (Poduslo and Curran 1996). Bioavailability is a major issue with new methods of delivery being sought. Since Tetanus toxin C (TTC) is effectively retrogradely transported to motor neurons following intramuscular delivery (Moreno-Igoa et al. 2010) it seemed that fusion proteins containing TTC may improve delivery of neurotrophins including BDNF to motor neurons. However, TTC and BDNF share the same internalization pathways and localize in the same endocytic compartments, suggesting that addition of TTC to the neurotrophins does not improve their bioavailability to motor neurons and may even prevent their entry (Roux et al. 2006). Indeed, when plasmids expressing recombinant fusion proteins containing BDNF and TTC were given to SOD1G93A mice there was no improvement of motor neuron health or survival above that of TTC (Calvo et al. 2011), suggesting that TTC competes more effectively than BDNF for entry into motor neurons. Lessons can also be learned from the signalling that is generated from activation of the TrkB receptor and its truncated form, TrkB.T1 One study showed that TrkB phosphorylation in ALS patients is decreased although the number of receptors is increased (Mutoh et al. 2000) suggesting a mechanism affecting the TrkB response to BDNF. In addition, knocking out the TrkB variant receptor TrkB.T1 receptors in SOD1G93A mice delayed the onset of motor neuron loss and functional symptoms (Yanpallewar et al. 2012) suggesting physiological or pathological imbalances in TrkB receptor isoform levels can also affect motoneuron survival.

### 4.4 Neurotrophin-3

Neurotrophin-3 (NT-3) is important for maintaining the integrity of neuromuscular junctions. It is well described that NT-3 increases retrograde transport within

degenerating motor neurons (Sagot et al. 1998) and enhances synaptic transmission (Poo 2001), in addition to stimulating neuronal survival. Importantly, muscles of NT-3-knockout mice lose neuromuscular innervation postnatally (Woolley et al. 2005) reflecting the requirement of peripherally derived NT-3 for neuron survival (Meier et al. 1999; Riethmacher et al. 1997) as well as for the maintenance of the neuromuscular junction (Reddy et al. 2003). NT-3 has not been trialled as a potential treatment. Importantly, it should be noted that NT-3, like BDNF, has an exceptionally short half-life in the circulation of only 1.28 min (Poduslo and Curran 1996).

### 4.5 Ciliary Neurotrophic Factor

The history of CNTF in ALS is similar to that of BDNF. CNTF was one of the earliest neurotrophic factors to be shown to have a direct neuroprotective effect on degenerating motor neurons in stress-induced conditions. Two studies in 1990 demonstrated that CNTF protects motor neurons in cell culture (Arakawa et al. 1990) and in vivo after axotomy-induced apoptosis in the rodent (Sendtner et al. 1990). Those results were strengthened by the beneficial effect of CNTF in a mouse model of neuronopathy (pmn) with motor neuron degeneration (Sendtner et al. 1992) and by increased vulnerability of spinal motor neurons in a knockout mouse line for CNTF (Masu et al. 1993). These results were published in the early 1990s, prior to the development of SOD1G93A mouse model of ALS (Gurney et al. 1994); today the most frequently employed model for ALS. Because of these promising effects on motor neurons, CNTF went directly to the clinic and became the first growth factor investigated clinically for ALS. Two clinical trials were conducted in 1996 and enrolled 570 and 730 patients, respectively (Miller et al. 1996; ALS CNTF Treatment Study Group 1996). The investigators focused mainly on daily living activities outcomes and survival. Unfortunately, both of those trials reported no observable benefit of treatment. Moreover, adverse effects, especially for the doses over 5  $\mu$ g/kg, were severe, leading to the cessation of treatment in many patients. What could be the reasons to explain the lack of efficiency of CNTF after such promising preclinical findings? Firstly, despite CNTF's ability to cross the BBB (Pan et al. 1999), a pharmacokinetic study in the rodent showed that CNTF has a half-life of only 3 min after intravenous injection (Poduslo and Curran 1996). Due to this low half-life, subcutaneous application of CNTF does not appear appropriate to target the CNS. Secondly, the doses used in animals were about 1 mg/kg, whereas the highest safe dose in the human is 200 times lower (5  $\mu$ g/kg). This limited dose and short half-life of CNTF could easily account for the lack of observable efficacy of CNTF in patients. No further clinical projects were pursued with CNTF in ALS after these two trials. A recent article from Sendtner's group (Selvaraj et al. 2012) has highlighted the promise of CNTF in stabilizing neuromuscular endplates and improving axon health. The action of a cytoplasmic protein called stathmin that destabilizes microtubules was inhibited by the activated signal transducer and activator of transcription-3 (STAT3). Importantly, STAT-3 was itself activated specifically by CNTF. This was done with motor neurons isolated from the pmn mouse model of neuronopathy; the same model used to show CNTF improves motor neuron health (Sendtner et al. 1992).

### 4.6 Insulin-Like Growth Factor-1

The action of IGF-1 in the nervous system (Shemer et al. 1987; Rotwein et al. 1988) was described long after its multitude of actions and targets in the body were known. Neurons in the central nervous system (CNS) express functional receptors of IGF-1 throughout the nervous system (Shemer et al. 1987; Rotwein et al. 1988; Bilak et al. 2001). IGF-1 is able to rescue motor neurons (Caroni 1993: Li et al. 1994) and reduce glutamate excitotoxicity (Nakao et al. 1996) both in vitro and in vivo. Therefore IGF-1 became an important drug candidate for ALS. Several clinical studies with IGF-1 in ALS were conducted early, before testing of IGF-1 in SOD1G93A rodents. In the late 1990s, two investigations were published with different conclusions (Lai et al. 1997; Borasio et al. 1998). Lai et al. enrolled 266 patients and administrated IGF-1 subcutaneously (Lai et al. 1997). The conclusion of the investigators was that IGF-1 is safe, slows progression of motor impairment, and increases quality of life. In a second trial, (Borasio et al. 1998) 183 patients were enrolled with the same mode of administration. The investigators focused on disease progression but found no improvement after treatment. Indeed, IGF-1 treatment appeared to be safe but did not improve quality of life of the patients nor motor functions. Additionally, a pilot study was conducted with intrathecal delivery of IGF-1 (Nagano et al. 2005b). Investigators claimed that the treatment was beneficial for the patients. Two groups of patients were compared, patients with a high and a low dose of IGF-1, and it was observed that the decline in motor function was slowed with the high dose of IGF-1, but not the low dose. However, as this study was not placebo controlled, the results should be considered with caution.

Many years after the first trials of IGF-1 in humans, preclinical reports investigating the effects of IGF-1 in SOD1 animals were published. The majority of the reports argue in favor of a potential protective effect in ALS, with an increase in survival and motor function parameters (Kaspar 2003; Nagano et al. 2005a; Lepore et al. 2007; Dodge et al. 2008; Franz et al. 2009). Also, IGF-1 mediates neuromuscular junction regeneration in a non-ALS-related animal model (Apel et al. 2010), although some contradictory results were published (Messi et al. 2007; Grumbles et al. 2009). For example, Messi et al. (2007) generated a transgenic form of SOD1G93A mice that overexpress IGF-1 in skeletal muscle but found no improvement of survival. The work of Franz et al. (2009) suggests that IGF-1 effects were only observed in male mice. At the same time, a new clinical trial was initiated with subcutaneous administration of IGF-1. This clinical trial enrolled 330 patients and studied the effects of IGF-1 on muscular strength and survival (Sorenson et al. 2008). Unfortunately, IGF-1-treated patients did not improve relative to the control patients.

There are a variety of reasons why IGF-1 failed at clinical trials. IGF-1 is increased at the spinal level of ALS patients (Wilczak et al. 2003) and in the

serum (Corbo et al. 2010). Conversely, muscle expression of IGF-1 and IGF-binding proteins was decreased in ALS patients (Lunetta et al. 2012), suggesting IGF-1 within the muscle tissue could participate in motor unit degeneration. However, as discussed above animal trials where IGF-1 is overexpressed in skeletal muscle in SOD1G93A mice did not improve survival (Messi et al. 2007). This may suggest the lack of success of the human clinical trial could be related to the transport of IGF-1 from muscle to the effected motor neurons and NMJ. In addition, it is hard to relate the preclinical trials involving viral gene delivery of IGF-1 (Kaspar 2003; Lepore et al. 2007; Dodge et al. 2008) or intrathecal administration (Nagano et al. 2005a) to the clinical trials where subcutaneous injections were used (Lai et al. 1997; Borasio et al. 1998; Sorenson et al. 2008). A genetic approach cannot be translated into subcutaneous delivery in patients, due to the high and constant expression of IGF-1 over time. Nor can the intrathecal administration that delivers IGF-1 to the CNS be directly comparable to subcutaneous injections. Indeed, IGF-1 has a half-life of only 4 min in the circulation (Pan and Kastin 2000), and intravenous IGF-1 enters the CNS by a saturable transport system at the BBB, which functions in synchrony with IGF-binding proteins in the periphery to regulate the availability of IGF-1. Therefore subcutaneous IGF-1 would seem unlikely to be an adequate mode of delivery. As Howe et al. (2009) observed it is highly unlikely that meaningful levels of IGF-1 were reached in the CSF and CNS tissue with the route and dose of IGF-1 used in the most recent clinical trial (Sorenson et al. 2008).

# 4.7 Vascular Endothelial Growth Factor

Discovered in the late 1980s as a growth factor for vascular endothelial cells, vascular endothelial growth factor (VEGF) induces endothelial cell growth and angiogenesis (Connolly et al. 1989). The first evidence for neurotrophic proprieties of VEGF arose with studies on neuronal regeneration (Sondell et al. 2000; Hobson et al. 2000). Shortly after, VEGF was implicated as a growth factor possibly involved in ALS by the discovery that mice with deletion of the HIF-response element in the VEGF promoter developed an ALS-like phenotype (Oosthuyse et al. 2001). Subsequent genetic analysis however failed to produce clear evidence for an effect of VEGF polymorphisms on the risk of developing ALS in humans (Gros-Louis et al. 2003; Lambrechts et al. 2003; Fernandez-Santiago et al. 2006; Chen et al. 2006; Zhang et al. 2006). A combined meta-analysis of American and European data showed the 2578AA VEGF genotype as a candidate for male ALS (Lambrechts et al. 2009). However, independent of the genetic discussion, VEGF promotes survival of cells overexpressing a mutated SOD1G93A protein (Li et al. 2003) and ameliorates the effects of glutamate excitotoxicity (Tolosa et al. 2008; Bogaert et al. 2010). In SOD1G93A rodents, VEGF treatment leads to a delay of disease onset, an improvement of motor functions, protection of motor neurons and neuromuscular junctions, and increase in survival (Zheng et al. 2004; Azzouz et al. 2004; Storkebaum et al. 2005; Wang et al. 2007; Hwang et al. 2009).

The preclinical data supports further clinical investigation of VEGF as a treatment target for ALS. Besides the animal studies, clinical observations show that VEGF is downregulated in the late stage of ALS (Devos et al. 2004), and restoring its level therefore could bring benefit for patients. The potential of VEGF as a therapeutic agent is further strengthened by its ability to cross the BBB (Wang et al. 1996; Dobrogowska et al. 1998; Ay et al. 2008). VEGF has an effective half-life of 10.7 min in the circulation (Wu et al. 2009). There are no published data on VEGF treatments for human ALS so far, but a clinical trial involving intracereb-roventricular infusion of VEGF (NCT0080050) with 28 patients is now at phase I/II. Another trial using plasmid-encoding zinc finger protein transcription factor that upregulates VEGF-A expression (NCT00748501) has completed phase II testing involving the treatment of 40 patients, but no outcomes have yet been published. Clearly, the angiogenic side effects of VEGF treatment will have to be monitored closely.

# 4.8 Fibroblast Growth Factors

The fibroblast growth factors (FGF) proteins are growth factors with homologous structure that promote angiogenesis and the proliferation of fibroblasts. It was demonstrated that both FGF-1 and FGF-2 can protect motor neurons after axotomy-induced apoptosis and that they support neuronal regeneration (Cuevas et al. 1995; Piehl et al. 1995; Grothe et al. 1991; Klimaschewski et al. 1999). No disease-specific alterations in the expression pattern of FGF-2 were found in postmortem spinal cord of ALS patients (Petri et al. 2009), but increased FGF-2 levels were measured in serum and cerebrospinal fluid of ALS patients (Johansson et al. 2003). Dependent on activation of different receptors, FGF-1 and FGF-2 can either promote cell survival or death (Grothe et al. 2006; Cassina et al. 2005). One report points out that FGF-1 might activate motor neuron apoptosis by astrocytes, thereby possibly contributing to ALS pathophysiology (Cassina et al. 2005). Subcutaneous administration of FGF-2 had no effect on the survival in transgenic SOD1G93A mice (Upton-Rice et al. 1999), but this may reflect the instability of native FGFs without heparin protection. A recent study where SOD1G93A mice were produced lacking endogenous FGF-2 showed increased lifespan of mice and upregulation of GDNF and CNTF (Thau et al. 2012). This supports the idea that endogenous FGF-2 contributes to the pathophysiology of SOD1G93A mice. Clearly, more work is needed to establish if the FGFs can be considered as drug candidates for ALS.

### 4.9 Hepatocyte Growth Factor

Hepatocyte growth factor (HGF) was discovered as growth factor acting on the liver (Nakamura et al. 1985). This growth factor exerts antiapoptotic activity in the liver after endotoxin-induced hepatic failure (Kosai et al. 1999). Neuronal effects consist of activity in the guidance and survival of neurons in the developing nervous

system (Ebens et al. 1996). Moreover, HGF is protective on motor neurons in culture and in vivo after axotomy (Wong et al. 1997; Okura et al. 1999). In rodent models of ALS, HGF reduces motor neuron degeneration and increases survival (Sun et al. 2002; Ishigaki et al. 2007; Kadoyama et al. 2007).

While HGF appears to be a good candidate for the treatment of ALS, especially since HGF levels in ALS patients appear to be dysregulated (Tsuboi et al. 2002), additional preclinical work is required. Indeed, more insight at the level of the NMJ is needed since it seems that HGF can have an influence on the formation of NMJs (Madhavan and Peng 2006; Tatsumi et al. 2009) but appears to be inefficient on their regeneration unless used in conjunction with other neurotrophic factors (Grumbles et al. 2009; Casella et al. 2010). There is work suggesting this may be because of pool-specific responsiveness of motor neurons to different neurotrophic factors. When the HGF receptor Met is turned off in mid-embryonic life, there is poolspecific loss of motor neurons at the NMJ within the pectoralis muscle from lack of Met (Lamballe et al. 2011). It will also be of interest for the design of a clinical trial to investigate whether a systemic delivery of HGF can reproduce the promising results observed after intrathecal administration in rodents (Ishigaki et al. 2007). HGF prevents the decline of choline acetyltransferase (ChAT) expression after adult hypoglossal nerve axotomy and also reduces motor neuron death in response to excitotoxic stress (Okura et al. 1999). Delayed disease onset and improved lifespan is observed when SOD1G93A mice overexpress transgenic HGF (Sun et al. 2002; Kadoyama et al. 2007). Reduced levels of the HGF receptor, Met, are found in the spinal cord of familial SOD1 patients over the course of disease, indicating disruption of the HGF-Met system contributes to neuronal degeneration in familial ALS (Kato et al. 2003). A recent study showed motor neuron development and maintenance are unaffected in mice with conditional enhancement of neuronal Met. whereas SOD1G93GA mice with the same enhancement of neuronal Met had a delayed disease onset (Genestine et al. 2011). Intrathecal HGF extends the lifespan of SOD1G93A mice, overcoming the HGF-Met deficit (Ishigaki et al. 2007). Another study showed in mice that the half-life of HGF is 10 min in the circulation, with 30 % of HGF crossing the BBB (Pan et al. 2006). A phase 1 clinical trial was intended to assess the safety and pharmacokinetics of single and multiple doses of rhHGF intrathecally in patients with ALS at Tohoku University Hospital (Japan) but was delayed because of the 2011 tsunami.

### 4.10 The Future of Neurotrophic Therapy for ALS

The demonstration of beneficial effects of neurotrophic factors for various CNS disease conditions raised great enthusiasm about their use as therapeutic agents for ALS. The clinical failure of neurotrophic factors is in no way ascribable to one single reason, especially in ALS.

Because of the failure to find clinical efficacy so far, the value of preclinical data has been questioned. There are two aspects to this issue. First, the models used may not be predictive for the working of a drug in the human system. This is of course an irrefutable argument, which cannot be proven or disproven in the absence of any clinical positive outcome seen so far. Second, animal models may be predictive but have not been conducted with sufficient rigor and diligence to really tell us the truth about efficacy of a drug candidate in the rodent system. Reasons for this include insufficient blinding procedures, neglect of copy number issues in transgenic models, insufficient number of animals used, inadequate statistical methods, and inadequate independent replication of study results (Scott et al. 2008). An example may have been BDNF, where preclinical studies were not done in SOD1G93A rodent models of ALS as they were not available at the time but were achieved using axotomy models, pmn and wobbler mice. After reports were published on the failure of the large clinical trial of BDNF (The BDNF Study Group 1999), there have been no effective BDNF-based treatments reported for the SOD1G93A ALS rodent models but one failure (Park et al. 2009).

Since there is insufficient proof that any one ALS animal model produced so far can be used to predict the effectiveness of human treatments, a combination of different animal models could be trialled. This could include different animal models that monitor different aspects of ALS disease pathophysiology, for example, two different SOD1-mutant models, possibly mouse and rat models, but with rodents that have adequate checks on copy number and preclinical trials using carefully designed studies. Ideally, a dose-effect relationship should be established. All tests have to be conducted in a rigorously blinded and randomized manner and ideally be performed in at least two independent laboratories. If novel animal models become available, they should be immediately tested using a variety of preclinical approaches.

In all major clinical trials performed so far, it appears the dosing was inadequate. Examples of this include BDNF, CNTF, and IGF-1 (The BDNF Study Group 1999; Ochs et al. 2000; Miller et al. 1996; ALS CNTF Treatment Study Group 1996; Lai et al. 1997; Borasio et al. 1998). For CNTF, the dose showing benefit for motor neurons in vivo was three orders of magnitude lower in the human case. In addition, subcutaneous-injected CNTF has a plasma half-life time of only a few minutes (Poduslo and Curran 1996) and does not reach the target tissues. A similar scenario can be drawn for BDNF, with questionable delivery to the CNS. For IGF-1, CSF levels reached with the chosen subcutaneous dose of 0.05 mg/kg twice daily, more than an order of magnitude below endogenous levels in the CSF, and thus are likely to be therapeutically meaningless (Howe et al. 2009).

The relevance of neurotrophic factor treatment for ALS should be pursued with a thorough understanding of how the neurotrophic factors work and if they address mechanisms relevant to ALS. For example, neurotrophic factors should show some efficacy against cell death either or both apoptotic and excitotoxic cell death demonstrated on motor neurons, other primary neurons, and/or motor neuron lines. Also, effects on the integrity of the NMJ should be ascertained. Only VEGF is known to preserve innervation to the NMJs in ALS (Zheng et al. 2007), although IGF-1 has shown benefit in preserving the NMJ after nerve transection (Apel et al. 2010). Addressing additive or synergistic effects of neurotrophic factors on NMJ as well as axonal and cell body health is an area yet to be adequately

explored as a treatment for ALS. However, there is an indication that some neurotrophic factors in combination improve NMJs; for example, cells transplanted with a combination of IGF-1, HGF, and GDNF appear to improve NMJs after transection, the combinatorial approach being better than individual factors (Grumbles et al. 2009; Casella et al. 2010). A recent article supports the notion of growth factor combinations having positive synergistic effects (Krakora et al. 2013). Mesenchymal stem cells were implanted into three muscle groups of SOD1G93A rats and secreted either GDNF, VEGF, IGF-1, or BDNF and compared to those implanted with cells secreting a combination of GDNF and VEGF (Krakora et al. 2013). The results indicated that rats treated with cells secreting both VEGF and GDNF have improved NMJ health and extended survival compared to rats treated with cells secreting individual neurotrophic factors. This indicates the need to further understand the cross talk between neurotrophic factors to improve their therapeutic effect in ALS.

For a chronic CNS disease like ALS, the application and dose chosen should guarantee a therapeutic level of the drug in the brain and spinal cord. For a motor neuron-supportive principle like neurotrophic factor support, the drug level should likely be held constant over time. Options for this include gene-based delivery of recombinant protein, including viral and nonviral methods, stem cell-based treatments that include secretion of relevant neurotrophic factors, or constant infusion pumps.

There is a good rationale for using other methods of delivery besides administration of the naked molecules, including half-life, getting to the site of action and prolonged neurotrophic factor production. For example, virally delivered genes for VEGF and IGF-1 have all shown promise in SOD1G93A rodents (Azzouz et al. 2004; Kaspar 2003; Lepore et al. 2007; Dodge et al. 2008). There is also a strong rationale for using nonviral delivery methods of targeting neurotrophic factors to specific sites of action such as motor neurons, NMJs, and glia. A nonviral gene delivery treatment capable of generating small quantities of therapeutic protein such as GDNF or HGF over a limited time targeting specific cells could have significant therapeutic value (Rogers and Rush 2012). This has been demonstrated for GDNF in an axotomy model in rats (Barati et al. 2006).

Stem cell-based delivery of neurotrophic factors has been suggested as an alternative method of delivering neurotrophic factors for ALS. Several studies report stem cell production of neurotrophic factors when transplanted into SOD1G93A rodents, including VEGF, IGF-1, GDNF, and BDNF (Hwang et al. 2009; Suzuki et al. 2007; Park et al. 2009). Cortical human neural progenitor cells engineered to secrete GDNF confer motor neuron protection after transplantation into the spinal cords of SOD1G93A transgenic rats (Suzuki et al. 2007). Intramuscular delivery of GDNF or GDNF-/VEGF-producing mesenchymal stem cells in SOD1G93A transgenic rats also increases neuromuscular contacts, motor neuron survival, and lifespan (Suzuki et al. 2008; Krakora et al. 2013). Coadministration of embryonic spinal cord cells with GDNF, HGF, and IGF-1 after sciatic nerve resection resulted in increased number of reinnervated muscle fibers and improved

motor neuron health (Grumbles et al. 2009). Human spinal stem cells (HSSC) that differentiate into both neurons and glia have been transplanted into SOD1G93A rodents and shown to secrete endogenous neurotrophic factors that may improve functional outcome. Following transplantation, HSSC express excitatory amino acid transporters that potentially restore functional glutamate reuptake around vulnerable motor neurons and also release neurotrophic factors such as GDNF and BDNF. Grafted HSSC express several neurotrophic factors, including GDNF and BDNF, and form synaptic contacts with host motor neurons (Xu et al. 2009). Moreover, intraspinal HSSC transplantation delays symptom onset and extends the lifespan in SOD1G93A rodents (Yan et al. 2006; Xu et al. 2006, 2011). Some caution has to be included as a recent study (Koh et al. 2012) has indicated that bone marrow mesenchymal stromal cells (BM-MSC) from ALS patients have a lower capacity to proliferate and secrete neurotrophic factors, indicating using autologous stem cells may be not be an effective in ALS patients.

# 5 Conclusion

It is nearly 20 years since the first trial of neurotrophic factors was reported for ALS. Since this time we have not only made significant inroads into understanding the pathophysiology and genetic cause of the disease, but we have gained more knowledge on the potential action of neurotrophic factors in ALS and proposed potentially smarter ways to deliver effective therapy. This provides reason for cautious optimism that a truly effective treatment is on the horizon.

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# Neurotrophins and p75NTR in Axonal Regeneration and Myelination

# Matt S. Ramer, Simon A. Bedard, and Angela L. M. Scott

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### Abstract

The neurotrophins are powerful, secreted signaling molecules that regulate the survival, proliferation, and function of neural cells. Neurotrophin-dependent signaling is not only critical during neurodevelopment but also plays an important role in the adult, where it mediates processes of plasticity and repair in both the central and peripheral nervous systems. The roles that the four known neurotrophins and their receptors play in regulating the cellular processes required for repair has been the subject of a significant body of research and has produced several promising avenues for treatment. This chapter reviews the current understanding of Neurotrophin-dependent signaling and how this evolving understanding can facilitate the development of future strategies to address

M.S. Ramer (🖂) • S.A. Bedard • A.L.M. Scott

International Collaboration on Repair Discoveries, The University of British Columbia, Vancouver, BC, Canada e-mail: ramer@icord.org

neurodegeneration and neurotrauma. Of particular interest in this context is the p75 pan-neurotrophin receptor, a mercurial molecule that binds not only the neurotrophins but also the other neurotrophin receptors in order to modulate these signaling cascades. P75-dependant signaling has been shown to be essential in the regulation of processes such as neurite outgrowth, axonal regeneration, cell survival, cellular proliferation, and myelination; these processes are critical to successful neural repair. As we learn more about the neurotrophins and p75 through future research, novel therapeutic targets for the treatment of damage to the nervous system will continue to be revealed.

#### Keywords

Brain-derived neurotrophic factor • Dorsal root entry zone • Dorsal root ganglion • Eph/ephrin • LINGO-1 • Myelination • Nerve growth factor • Nerve growth factor receptor • Neurotrophin-3 • NogoA receptor • Oligodendrocyte • Semaphorin • Sortilin • Trk receptor

### 1 Introduction

Since the discovery of nerve growth factor (Cohen et al. 1954), neurotrophins (NTs) have been touted as potential remedies for neurotrauma. For injuries involving peripheral nerves or the spinal cord, the problem is inefficient or ineffective growth of damaged axons. The promise of NTs stems from the notion that disability may be reversed by providing neurons and their associated glia with adequate trophic support. While much work has been aimed at exploiting the function of spared axons as a possible solution to incomplete neurotrauma (Ramer 2010), true functional recovery entails regeneration of damaged axons and their reconnection with appropriate targets. This is not all, however, since regenerated axons must be myelinated to reliably conduct action potentials. It is therefore not insignificant that NTs provoke multiple responses from myelinating glia.

In this chapter the biology of NTs and their receptors will be reviewed, with a particular emphasis on the pan-NT receptor p75NTR, given its pivotal role in a growing number of signalling mechanisms. The effects of NTs and their receptors on regeneration and myelination, mainly in the context of sensory neurons of the dorsal root ganglion (DRG), will be discussed. By virtue of their anatomical arrangement (with axons in both the peripheral and central nervous systems) and well-characterized responses to NTs, DRG neurons have proven particularly informative with respect to the mechanisms by which NTs mediate both regeneration and myelination.

# 2 The Neurotrophins

Neurotrophic factors are characterized by their ability to support the survival and growth of specific neuronal populations. The neurotrophic hypothesis holds that

target-derived neurotrophins, retrogradely transported to cell bodies, are responsible for maintenance of neuronal phenotype in adulthood. The most complete existing data relates to the NTs (Bradbury et al. 2000; Boucher et al. 2000). There are four known NTs: nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), neurotrophin-3 (NT-3), and neurotrophin-4 (NT-4), all of which form homodimers (Chao 2003). Each NT is derived from a precursor molecule (pro-NT) that can be cleaved by a furan or convertase to produce a mature NT which is subsequently released into the extracellular space. Although this posttranslational modification occurs intracellularly, unprocessed pro-NTs are also released to the extracellular space via the constitutive secretory pathway and – in the case of pro-BDNF – via a regulated secretory pathway (Mowla et al. 2001). The mature forms are non-covalently bound dimers that share 90 % homology, including a cysteine-rich motif within the structural core of the dimer and three beta hairpin loops forming the distinctive regions involved in determining receptor specificity (Wiesmann and de Vos 2001).

# **3** Neurotrophin Receptors

### 3.1 Trk Receptors

NT signals are received and transduced by the tropomyosin-related kinase (Trk) A, B, and C receptors as well as the p75 neurotrophin receptor (p75NTR), the latter being a member of the tumor necrosis factor receptor (TNFR) family. Each Trk receptor is a cell surface transmembrane protein that possesses ligand-dependant tyrosine kinase activity. However, alternative splicing of TrkB (TrkBT1, TrkBT2, and TrkBT4) and TrkC (TrkCTK<sup>-</sup>) produces additional truncated isoforms of these receptors that lack the tyrosine kinase domain (Squinto et al. 1991; Valenzuela et al. 1993; Forooghian et al. 2001). Each full-length receptor and truncated isoform share a conserved extracellular domain that includes two immunoglobulin-like C2 domains that not only prevent the autoactivation of the full-length receptors (Arevalo et al. 2000) but also determine neurotrophin binding specificity (Urfer et al. 1995, 1997). This specificity has translated into high-affinity interactions of NGF with TrkA, BDNF and NT-4/5 with TrkB isoforms, and NT-3 with TrkC isoforms (Chao 2003). Thus in the majority of cases, the cellular sensitivity and responsiveness to neurotrophins is dictated by the expression of their respective Trk receptors (although NT-bound p75NTR can participate in NT signalling through alternate pathways (Naska et al. 2010)).

# 3.2 p75NTR

Perhaps the most interesting member of the neurotrophin-signalling suite is p75NTR, whose many roles in cellular signalling continue to emerge. p75NTR was first identified in 1973 (Herrup and Shooter 1973) and was originally named the nerve growth factor receptor (NGFR); the name was later changed when it was

realized that p75NTR had affinity not only for NGF but also for the other NTs. p75NTR was subsequently characterized as a 75 kDa, type I transmembrane protein (Grob et al. 1985). p75NTR has been simplistically described as a regulator of cell survival or death, but in reality its biological function is much more subtle and nuanced. In fact, p75NTR signalling can have effects as wide reaching as cell death, proliferation, differentiation, survival, neurite outgrowth/collapse, and even the modulation of synaptic connectivity, plasticity, and pruning (Skeldal et al. 2011; Garcia et al. 2011).

### 3.3 p75NTR Signalling Partners

There are several signalling modules in which p75NTR participates. First, p75NTR has been shown to greatly increase the binding affinity of the Trk receptors for their cognate NTs (Esposito et al. 2001). The Trk receptors form receptor dimers upon ligand binding, as do most receptor tyrosine kinases, and trans-autophosphorylate one another to activate downstream signalling pathways (Chao 2003). The association of p75NTR with these Trk dimers creates high-affinity binding sites, increasing both the sensitivity and degree of Trk signalling. In this way p75NTR is capable of positively regulating Trk signalling. The converse, however, is also possible. With proteolytic cleavage at a juxtamembrane  $\alpha$ -secretase site, the extracellular domain of p75NTR is released into the extracellular space where it can bind competitively with available NTs. The consequent dearth of NTs available for the Trk receptors results in decreased Trk signalling.

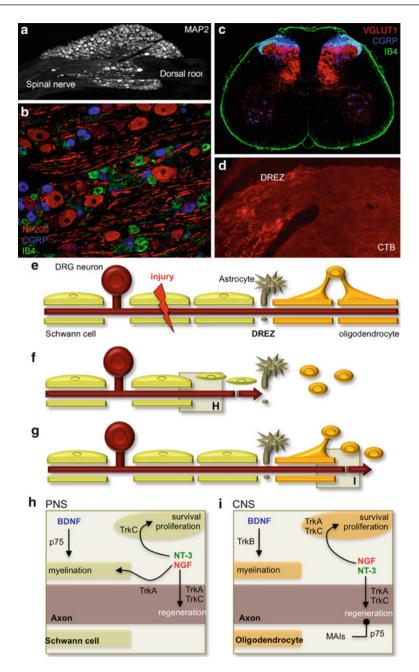
Depending on the NTs present, the Trk receptors being expressed, and the type of cleavage (if any) that p75NTR undergoes, a whole multitude of functional outcomes is possible. Each of the different Trk receptors activates different downstream signalling pathways, which are often dependent on p75NTR. For example, TrkA is not able to activate a PI3-kinase/Akt signalling cascade without associating with p75NTR, but can activate a MAP-kinase cascade with or without p75NTR (Nykjaer et al. 2005). The Trk receptors are broadly involved in cell proliferation, differentiation, and survival, and p75NTR serves to enhance or mute these effects when it does or does not associate with them.

The Trk receptors are not p75NTR's only bedfellows, however. p75NTR also forms a complex with sortilin, a protein usually involved in intracellular vesicle trafficking, at the plasma membrane (Nykjaer et al. 2004). The association of sortilin with p75NTR greatly increases the affinity of p75NTR for the pro-NTs over mature NTs. It was shown that the interaction between p75NTR and sortilin is necessary for pro-NGF- and pro-BDNF-induced apoptosis (Nykjaer et al. 2004). In this context p75NTR is no longer a peripheral player, serving to modulate the signalling of another receptor, but is rather the receptor that transduces the signal to the cytosol. Upon binding of a pro-NT by the p75NTR/sortilin complex, the intracellular domain of p75NTR is cleaved and induces apoptosis via a Jun N-terminal kinase (JNK)-dependent mechanism (Barker 2004; Kenchappa et al. 2010).

A third signalling module that p75NTR is involved in is a membrane-bound complex with *LINGO-1* and, in neurons, the NogoA receptor (NgR) (Mi et al. 2004). This complex does not signal in response to NT or pro-NT signals, but rather responds to myelin-associated inhibitors (MAIs). These include Nogo, myelin-associated glycoprotein (MAG), and oligodendrocyte myelin glycoprotein (OMgp) (Wang et al. 2002). In this circumstance p75NTR (or the related molecule Troy) (Park et al. 2005; Shao et al. 2005) is again required for signalling. Although LINGO-1 possesses a tyrosine kinase phosphorylation site in its cytoplasmic aspect – indicating that it is likely able to interact with downstream intracellular signalling cascades (Mi et al. 2008) – it is unable to induce signalling in the absence of p75NTR, despite being able to bind its ligands. In neurons, the functional effect of signalling through the p75NTR/NgR/LINGO-1 complex is inhibition of neurite outgrowth via collapse of extending axons' growth cones (Yamashita and Tohyama 2003). In recent work, Park et al. (Park et al. 2010) showed that developmental MAI-dependent axonal pruning is also p75NTR dependent.

p75NTR is also connected to a fourth receptor complex, involved in Eph/ephrin signalling. In 2001, Kong et al. (Kong et al. 2001) cloned a novel transmembrane protein they called ARMS (ankyrin repeat-rich membrane spanning) downstream of Eph and neurotrophin signalling. ARMS interacted with both TrkA and p75NTR and could be phosphorylated following NT or ephrin B treatment of cultured cells. This was an exciting result, particularly since ephrin B3 was subsequently shown to be another MAI (Benson et al. 2005): p75NTR/ephrin B signalling may participate in ephrin B3-mediated neurite growth inhibition. Bolstering this notion, in sympathetic neurons p75NTR is required for ephrin B-mediated growth cone collapse (Naska et al. 2010). Other recent and compelling evidence of inhibitory p75NTR/ephrin A "reverse" signalling has emerged (Lim et al. 2008). Ephrin A lacks a transmembrane domain, but in the retinotectal system, it couples with p75NTR to mediate retinal axon repulsion during topographic map formation.

Inhibitory semaphorin signalling in growing axons also seems to involve p75NTR, although the data are not as clear-cut. p75NTR has been reported to bind plexins and neuropilins, both receptor components for semaphorin family members (Ben Zvi et al. 2007, but see Naska et al. 2010). In the absence of p75NTR, sensory neurons are hypersensitive to Sema 3A, an effect thought to be the result of enhanced association of neuropilins with plexins. Hypersensitivity of p75NTR sympathetic axons to Sema 3A was also shown in p75NTR-/- hearts, where the high-to-low gradient of sympathetic innervation from epi- to endocardium – shaped by endocardial Sema 3A expression during development (Ieda et al. 2007) - was greatly exaggerated (Lorentz et al. 2010). This same study also showed increased Sema 3A-mediated inhibition of outgrowth from p75NTR-/- sympathetic ganglion explants. These findings stand in contrast to another showing that functional p75NTR, bound to a NT ligand, is actually required for the growth cone collapsing effects of Sema 3 family members on sympathetic neurons (Naska et al. 2010). Although it seems clear that p75NTR participates in semaphorin signalling, there is not yet a complete picture of how it does so.



**Fig. 1** Anatomy and selected NT responses of sensory neurons and associated glia. (a) A lumbar DRG section from a rat. (b) Subtypes of DRG neurons. Blue and green neurons, nociceptors; red neurons, mechano-/proprioceptors. (c) Central projections of sensory neurons. Terminal projections of subpopulations correspond to colors used in **B** (pia mater also binds IB4, and some motoneurons also express CGRP). (d) Large-diameter DRG neurons injured in the dorsal roots 1

### 4 Neurotrophins and Sensory Axon Regeneration

The tenets of the neurotrophic hypothesis have been most extensively demonstrated for sensory neurons of the dorsal root ganglion (DRG) (McMahon et al. 1995; Verge et al. 1995; Munson et al. 1997). As such, DRG neurons are ideal for assessing the role of neurotrophic factors and their receptors in regeneration. The anatomical arrangement of DRG neurons makes them ideal for experimental manipulation: they are pseudounipolar in that each has a single axon which emanates from the soma and bifurcates within the DRG, sending one branch peripherally in the peripheral nerves and the other centrally in the dorsal roots (Fig. 1). The dorsal root axons pass from the PNS into the CNS at a well-defined border, the dorsal root entry zone (DREZ). Once beyond the DREZ, most axons bifurcate again, forming longitudinal bundles. These are the ascending dorsal columns, large myelinated axons which terminate in the brainstem, and Lissauer's tract, a primarily unmyelinated propriospinal tract. From the longitudinally projecting axons extend branches which terminate in the grey matter of the dorsal and ventral horns. These are clustered around the axon's point of entry into the cord, but can occupy several segments rostrally and caudally. Thus, sensory axons can be injured in the peripheral nerve, in the dorsal roots (by rhizotomy), or within the spinal cord (Bradbury et al. 2000).

Sensory neurons come in several phenotypes, and this heterogeneity has been investigated in exquisite detail. About two thirds of all DRG neurons have unmyelinated or thinly myelinated axons, and more than 90 % of these innervate nociceptors peripherally. Roughly half of these are "peptidergic," as they express peptide neurotransmitters substance P and calcitonin gene-related peptide (CGRP). The remaining third of DRG neurons have rapidly conducting myelinated axons and subserve mechanoreception and proprioception.

All of the receptor components for neurotrophins are expressed in the adult DRG. Interestingly, small, substance P and CGRP-expressing nociceptive neurons virtually all express NGF receptor components – TrkA and p75NTR. The non-peptidergic group expresses none of the full-length Trk receptors, nor p75NTR (Molliver et al. 1997; Bennett et al. 1998). More than half of the large light neurons express TrkC and p75NTR, while few adult DRG neurons express TrkB (McMahon et al. 1994). Thus, the majority of sensory neurons respond to either NGF or NT-3.

NTs promote outgrowth of the expected phenotypes of adult sensory neurons (based on receptor expression in vivo) in dissociated culture (Lindsay 1988;

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**Fig. 1** (continued) week earlier and halted at the DREZ. (**e**–**g**) Stages in successful regeneration across the DREZ following dorsal root injury (indicated in **G**). (**h**, **i**) NT responses of axons, Schwann cells, and oligodendrocytes (and their precursors) and the NT receptors involved. Abbreviations: MAP2, microtubule-associated protein 2; NF200, 200 kDa neurofilament; CGRP, calcitonin gene-related peptide; IB4, lectin from *Bandeiraea simplicifolia*; VGLUT1, vesicular glutamate transporter 1; CTB, cholera toxin B fragment; MAIs, myelin-associated inhibitors

Hu-Tsai et al. 1994; Tuttle and Matthew 1995; Kimpinski et al. 1997; Gavazzi et al. 1999): NGF and NT-3 enhance outgrowth from peptidergic nociceptors and NF200-positive mechano-/proprioceptors, respectively. BDNF seems to have varied effects on sensory neurons in vitro, in some cases promoting outgrowth and in others inhibiting it (Gavazzi et al. 1999). In vivo, BDNF is ineffective at promoting regeneration of large-diameter axons (Bradbury et al. 1999; Ramer et al. 2000); furthermore, it inhibits intraspinal collateral sprouting of most primary afferent phenotypes (Ramer et al. 2007; Soril et al. 2008).

While the in vitro data show that NTs are clearly sufficient to elicit neurite outgrowth, there appears to be little necessity for endogenous NTFs in naturally occurring regeneration of peripheral nerves. For example, axotomized TrkA- and TrkC-expressing nociceptive and mechanosensitive axons reinnervate their peripheral targets normally in the presence of antibodies which block NGF and NT-3 signalling (Diamond et al. 1987; Yasargil et al. 1988; Diamond et al. 1992). Surprisingly little is known about the efficacy of exogenous NTs in enhancing regeneration of sensory axons following peripheral nerve injury. This is possibly because peripheral nerve regeneration actually happens, making subtle increases in regeneration distance or numbers of regenerating axons difficult to detect.

The dorsal rhizotomy model is particularly attractive for studying regeneration by virtue of the fact that regeneration proceeds in the permissive Schwann cell environment of the injured dorsal root, but halts upon contact with the astrocytic glia limitans at the PNS-CNS interface. Thus the role of neurotrophins and their receptors can be examined on a background of unequivocal regeneration failure. Using this model, Ramer et al. (Ramer et al. 2000) found that intrathecal delivery of NGF and NT-3, but not BDNF, resulted in regeneration of injured sensory axons beyond the DREZ and into the spinal cord. As might be expected from prior in vitro work, the phenotype of regenerating axons corresponded to Trk expression: NGF resulted in ingrowth of CGRP-expressing (TrkA-positive) primary afferents, and NT-3 resulted in regeneration of (TrkC-expressing) NF200-positive axons. The axons which regenerated were able to synaptically drive second-order neurons (Ramer et al. 2000, 2002), and at least in the case of NT-3, there was ultrastructural evidence of new synapses with neurons in the dorsal grey matter (Ramer et al. 2002). NGF administration was accompanied by the reemergence of responses to heat and noxious pressure (Ramer et al. 2000), and in NT-3-treated animals, behaviors requiring proprioception such as paw placement and walking across a beam and across a horizontal ladder returned (Ramer et al. 2002).

### 5 p75NTR and NT-Mediated Sensory Axon Regeneration

An attractive model for p75NTR/Trk interactions in axonal regeneration was posited in the late 1980s (Johnson, Jr. et al. 1988). This involved a "presentation" mechanism, by which Schwann cell-expressed p75NTR generated an NT gradient which could be followed by axons. This hypothetical framework led to the widely accepted notion that the powerful chemotactic effects of NTs on axons are mediated

by p75NTR (Huang and Reichardt 2003). However, several studies have pointed to a negative role of p75NTR in NT signalling and neurite outgrowth. Most notably, regeneration of motor axons following peripheral nerve injury is more robust in p75NTR knockouts than in wild-type mice (Ferri et al. 1998; Boyd and Gordon 2001, but see Song et al. 2009).

These data, combined with the possibility that  $\alpha$ -secretase disrupts any substratebound gradient by liberating the ECD of p75NTR (Zampieri et al. 2005), led to questions regarding the validity of the presentation model. In fact, in p75NTR - / mice (exon III deletion), injured sensory axons were able to not only penetrate the dorsal root entry zone and enter the CNS following dorsal rhizotomy but also drive CNS neurons and restore sensory function (Scott and Ramer 2010). Successful regeneration was dependent on endogenous NGF and NT-3, as the regeneration of both small-caliber and large-diameter axons was prevented in p75NTR - /-miceby NGF- and NT-3 sequestration by TrkA-Fc and TrkC-Fc, respectively. Importantly, p75NTR-/- axons could not penetrate the astrocytic scar unless their associated Schwann cells were also p75NTR - / -, and in tissue culture experiments of Schwann cells and DRG neurons, it was shown that Schwann cell-expressed p75NTR reduced NT-dependent neurite outgrowth. Since NGF and NT-3 levels were equivalent in wild-type and p75NTR - /-mice, these data showed that rather than presenting neurotrophins to regenerating axons, Schwann cell p75NTR restricts NT availability and limits regeneration into the cord.

It is important to contrast this work with studies on the effects of p75NTR following injury to sensory axons within the CNS. Since p75NTR transduces inhibitory signalling by MAIs, it is reasonable to hypothesize that its absence might relieve inhibition. However, Song et al. (Song et al. 2004) showed that sensory axons injured in the dorsal columns of p75NTR - /-mice (the same exon III null mutants used in the rhizotomy studies) failed to regenerate beyond the glial scar. The crucial difference between this and the rhizotomy model is the lack of reactive and NT-producing Schwann cells following direct CNS lesions. That is, regeneration did not fail following dorsal column injury because p75NTR was unable to mediate MAI signalling, but because trophic support was inadequate. Indeed, grafts of p75NTR - / - peripheral nerve are better able to support growth of injured dorsal column axons than are wild-type grafts (Scott and Ramer 2010). Together, these studies cast doubt on the role of all known p75NTR-interacting inhibitory molecules produced in the injured spinal cord (including NogoA, MAG, OMgp, ephrin B3, and Semaphorin 3A) in blocking regeneration of injured axons. In fact, a recent study using mice lacking all three known NgR ligands showed a frank absence of regeneration, reinforcing these uncertainties (Lee et al. 2010).

### 6 Neurotrophins, p75NTR, and Myelination

Successful establishment of axonal conduction during development, and successful recovery of function following neurotrauma, requires myelination. Some of the most

promising experimental strategies for promoting axonal outgrowth following CNS injury include manipulating neurotrophic factor levels and antagonizing signalling by MAIs (although in the latter case "outgrowth" is restricted to plasticity of spared fibers rather than regeneration of injured axons (Lee et al. 2010)). Despite its importance to functional recovery, it has yet to be determined in any meaningful way whether myelination accompanies the axonal regeneration induced by these manipulations.

In the PNS, axonal internodes are myelinated by Schwann cells (One Schwann cell per internode), and in the central nervous system (CNS), this task is performed by oligodendrocytes (multiple internodes per oligodendrocyte). Although similar in many respects, there are some very marked differences between these two cell types beyond their location and structure. One such difference is the expression of p75NTR following injury (Zhou and Li 2007). In Schwann cells p75NTR is dramatically upregulated following injury and remains so until axon/Schwann cell contact is reestablished, while in oligodendrocytes this is not the case. It has been suggested throughout the literature that this difference may be one key aspect that prevents CNS damage from being repaired in the same way as damage to the PNS, although the proposed mechanisms by which this might be accomplished vary.

# 6.1 Schwann Cells

In Schwann cells, Trk signalling is responsible for regulating proliferation and differentiation, as well as elongation and axon ensheathment. Cosgaya et al. (2002) demonstrated that NT-3, via TrkC, has a powerful inhibitory effect on myelination, maintaining Schwann cells in a proliferative state. NT-3 signalling via p75NTR also negatively affects Schwann cell migration (Bentley and Lee 2000; Yamauchi et al. 2004; Skeldal et al. 2011). Once NT-3 levels begin to drop, however, BDNF is able to upregulate myelin-specific genes in Schwann cells. BDNF is central to myelination by Schwann cells both in development and after injury: BDNF derived from sensory axons promotes Schwann cell myelination during development (Tolwani et al. 2004; Ng et al. 2007), and, following peripheral nerve injury, BDNF is required for myelination of regenerating axons (Zhang et al. 2000). Crucially, in all cases BDNF exerts its effects on Schwann cells via p75NTR (Cosgaya et al. 2002; Song et al. 2006).

NGF also promotes myelination in DRG-Schwann cell cocultures (Chan et al. 2004), although this is mediated not by a direct action of NGF on Schwann cells, but via TrkA expressed on sensory axons. It was speculated that some of the effects might have been due to an increase in axonal diameter, but it is well known that NGF potently upregulates BDNF selectively in TrkA-expressing DRG neurons and their processes (Michael et al. 1997). The potent effects of BDNF on myelination by Schwann cells probably underpin these indirect effects of NGF.

Schwann cell p75NTR not only is involved in the initiation of the glial myelination program and the maintenance of cell survival but appears to be involved in the actual act of ensheathment as well. Chan and colleagues (Chan et al. 2006) demonstrated that p75NTR is actively recruited to the axon-glial junction by the

polarity protein Par-3 at the commencement of ensheathment. Moreover, they showed that axon-Schwann cell contact was necessary to induce this recruitment and localization of p75NTR and proposed that this positions p75NTR precisely on the plasma membrane of the Schwann cell to receive pro-myelinating signals.

## 6.2 Oligodendrocytes

Neurotrophin signalling also operates in the oligodendrocyte lineage. Functional Trk receptors are expressed on these cells (Barres et al. 1994; Condorelli et al. 1995; Cohen et al. 1996; Kumar and de Vellis 1996); NGF and NT-3 promote oligodendrocyte precursor cell (OPC) survival in vitro and, when combined with plateletderived growth factor, NT-3 induces their proliferation (Barres et al. 1994). NGF inhibits oligodendrocyte myelination of sensory axons in culture, but this occurs through modulating the "receptivity" of TrkA-expressing axons rather than via direct effects on OPCs (Chan et al. 2004). NT-3 withdrawal from culture results in OPC differentiation into pre-myelinating oligodendrocytes, and this is generally accepted to be the result of a cell-autonomous cycle-counting mechanism (Barres and Raff 1994; Durand and Raff 2000). In contrast to the survivaland proliferation-enhancing effects of NT-3, BDNF initiates the myelination program (via TrkB), and, accordingly, BDNF heterozygous mice exhibit delayed developmental myelination (Xiao et al. 2010). Unlike in Schwann cells, BDNF promotes oligodendrocyte myelination equally well in the presence or absence of p75NTR (Du et al. 2006; Xiao et al. 2010).

This is not to say that NGF and NT-3 never push oligodendrocytes toward differentiation. NT-3 and NGF have been shown to increase the expression of myelin basic protein and proteolipid protein, both components of mature myelin, in basal forebrain oligodendrocytes (Du et al. 2006). Unlike BDNF, they do so in part via p75NTR. For NT-3 at least, this occurs through activation of myelin protein translation, rather than by increasing myelin gene transcription (Coelho et al. 2009). Additionally, high concentrations of NT-3 are capable of transiently inducing OPC differentiation into oligodendrocytes (Heinrich et al. 1999).

Much of what is known about NT effects on oligodendrocytes comes from in vitro studies and developmental assessment of myelination in transgenic mouse studies; very little is known about NT effects on oligodendrocytes, their progenitors, or myelination of axons in the adult animal following neurotrauma. The dorsal rhizotomy model may again be particularly informative here, since Schwann cells do not invade the CNS when the injury occurs in the dorsal roots. In quantitative stereological TEM studies on rhizotomized rats, it was shown that 1 week's intrathecal NT-3 treatment, which promotes regeneration of sensory axons into the spinal cord, also results in a near doubling of oligodendroglial tissue in the CNS part of the DREZ (Hanna-Mitchell et al. 2008). How NT treatment affects myelination of regenerated axons in this model has not yet been worked out, but this task should be made easier by the fact that rhizotomy – unlike direct CNS trauma– does not result in oligodendrocyte progenitor death or demyelination of spared axons (Sun et al. 2010).

p75NTR/LINGO-1 signalling also has a role to play in oligodendrocytes, but it is distinct from that in neurons. In oligodendrocytes this signal inhibits differentiation of OPCs into pre-myelinating oligodendrocytes (Mi et al. 2005). LINGO-1 signalling via p75NTR seems to be essential to oligodendrocyte process outgrowth (and by extension, myelination), since the inhibitory effects of LINGO-1 are reversed by p75NTR antagonists (Bourikas et al. 2010). Intriguingly, axonal LINGO-1 also inhibits oligodendrocyte differentiation, an effect that is potentiated by NGF acting through axonal TrkA (Lee et al. 2007). LINGO-1 is expressed in white matter of the injured spinal cord (Barrette et al. 2007), so it is reasonable to speculate that myelin-derived inhibitory proteins released into the extracellular space following injury inhibit nearby OPCs from differentiating into myelinating oligodendrocytes. This intriguing hypothesis has yet to be tested, however.

# 7 Conclusion

The literature reviewed in this chapter illustrates the wide-ranging effects of NTs on axonal regeneration and myelination, both prerequisites for functional recovery in the wake of neurotrauma. These have largely been examined in isolation, however, and the time has come to develop strategies aimed at optimizing both processes. Although details vary, the following generalizations can be safely made (Fig. 1): NGF and NT-3 promote regeneration of sensory axons, and while NGF promotes peripheral myelination, NT-3 inhibits it. NGF and NT-3 also tend to enhance OPC survival and proliferation. BDNF, on the other hand, not only promotes myelination by both Schwann cells and oligodendrocytes but also inhibits plasticity of sensory axons. p75NTR is required for appropriate peripheral myelination, but may act to prevent oligodendrocyte myelination through its association with LINGO-1. For regeneration in the CNS – one of the toughest challenges in neuroscience – one might envision a sequential approach using NTs, first targeting sensory axonal regeneration and then initiating the myelination program.

The hope that neurotrophins hold for repairing the damaged nervous system must be tempered by the understanding that developmental mechanisms underlying neurotrophin-mediated axon growth and myelination are unlikely to be faithfully recapitulated in the adult. However, manipulations involving powerful experimental models – such as primary afferent neurons and their processes – will continue to shed light on how neurotrophin treatment might be refined to achieve all requirements of functional restoration.

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# **Neurotrophins and Pain**

# Ru-Ping Dai and Xin-Fu Zhou

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#### Abstract

There is ample evidence suggesting that neurotrophins in particular nerve growth factor (NGF) and brain-derived neurotrophic factor (BDNF) regulate pain processing as the modulators. NGF released from local tissue and dorsal root ganglia (DRG) binds to tropomyosin-related kinase (Trk) A receptors and

R.-P. Dai (🖂)

e-mail: Ruping\_dai@yahoo.com.cn

X.-F. Zhou

Department of Anesthesia, The Second Xiang-Ya Hospital of Central South University, Changsha, Hunan Province, China

School of Pharmacy and Medical Sciences, Faculty of Health Sciences, University of South Australia, Adelaide, SA, Australia e-mail: xin-fu.zhou@unisa.edu.au

regulates many membrane receptors, ion channels, and signaling molecules contributing to pain processing. Molecules antagonizing NGF-TrkA signaling have been developed and attenuate chronic pain in the patients. However, anti-NGF therapy leads to bone death in some patients suggesting that the therapy antagonizing NGF-TrkA signaling should be cautious to control pain in patients with chronic pain. On the other hand, brain-derived neurotrophic factor (BDNF) may regulate pain as a central modulator. BDNF is released when nociceptors are activated and regulated by NGF. This chapter reviews the roles and the underlying mechanisms by which NGF and BDNF regulate pain processing. The clinical trial of anti-NGF therapy is also discussed.

#### Keywords

Brain derived neutrophic factor • Central sensitization • Nerve growth factor • Neurotrophin • Pain • Spinal cord

#### List of Abbreviations

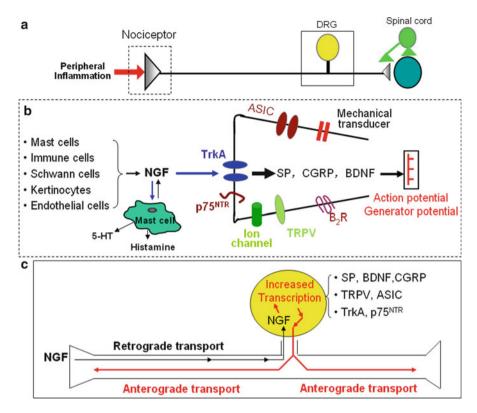
LIST OF ADDICTION	
5-HT receptor	5-Hydroxytryptamine receptor
ASIC	Acid-sensing ion channels
$B_2R$	Bradykinin (BK) receptors
BDNF	Brain-derived neurotrophic factor
CGRP	Calcitonin gene-related peptide
CREB	cAMP response element-binding protein
DRG	Dorsal root ganglia
EPSCs	Excitatory postsynaptic currents
ERK	Extracellular regulated kinase
GABA	γ-aminobutyric acid
NGF	Nerve growth factor
NMDA receptor	<i>N</i> -methyl-D-aspartate receptor
NSAID	Nonsteroidal anti-inflammatory drugs
NT	Neurotrophin factor
p75 <sup>NTR</sup>	p75 neurotrophin receptor
SP	Substance P
TNF-α	Tumor necrosis factor-α
TRK	Tropomyosin kinase
TRPV	Transient receptor potential vanilloid

## 1 Introduction

Pain is an unpleasant feeling often caused by noxious stimuli such as trauma, stubbing a toe, or burning the finger. As defined by the International Association for the Study of Pain, pain is "an unpleasant sensory and emotional experience associated with actual or potential tissue damage, or described in terms of such damage" (Bonica 1979). It can motivate the individual to withdraw from the damaged situation and is essential for the avoidance of tissue injury and promotes healing after injury (Lynn 1983). Pain is usually transitory and disappears once the noxious stimuli are removed or the injured tissue is healed. However, in many situations, pain can be persistent even after removal of the noxious stimuli or the healing of the body. The development of pathological or chronic pain will severely affect the patients' life quality and general functions (Raj 2007). As a matter of fact, pain is the most common reason for physician consultation in the United States and Europe, and management of pathological pain is still a major health challenge throughout the world (Breivik et al. 2006; Hardt et al. 2008). This is because there is still a lack of effective treatments of chronic pain without unwanted side effects and/or abusive liability.

Currently, nonsteroidal anti-inflammatory drugs (NSAID) such as ibuprofen and cyclooxygenase-2 inhibitors and opioids such as tramadol or morphine are most widely used for pain control in clinical practice. However, the concerns of gastro-intestinal and renal side effects of NSAIDs and of cardiovascular risk of cyclooxygenase-2 inhibitors limit their use in many medical conditions (Whelton 2000). On the other hand, despite the fact that opioids are an effective treatment in the many types of chronic pain, their side effects and liabilities, including loss of drug effectiveness, constipation (the most common long-term side effect causing noncompliance), drug diversion, respiratory depression and accidental death caused by overdose, are major concerns in clinical practice (Chou et al. 2009). Therefore, development of new agents for chronic pain control without severe side effects remains a significant, unmet clinical need.

Neurotrophins are a family of growth factors containing the following members: nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), neurotrophin factor (NT), and NT-4/5. All neurotrophins are synthesized as a proform of approximately 30 kDa and cleaved to a mature form of approximately 13 kDa (Barbacid 1995; Reichardt 2006). Each neurotrophin binds with high affinity to their specific receptors, one of the tropomyosin kinase (Trk) family of transmembrane receptors. For example, NGF binds to TrkA, BDNF and NT-3/4 bind to TrkB, and NT-3 binds to TrkC (and to TrkA with lower affinity). Conversely all neurotrophins have a common low-affinity binding transmembrane receptor, p75 neurotrophin receptor (p75<sup>NTR</sup>). Through binding to their highaffinity receptors and activating the downstream signaling, neurotrophins can regulate the survival, development, and function of neuronal subpopulations that express appropriate receptors (Barbacid 1995). In the absence of Trk or if the ratio of p75/TrkA is high, neurotrophins can activate p75<sup>NTR</sup> and subsequently the downstream signaling (JNK, Nf-kB) and promote apoptosis. In addition to regulating the survival and function of the nervous system, there is ample evidence indicating that NGF and BDNF play important roles in development of various types of pain (Pezet and McMahon 2006; Mantyh et al. 2011). In this chapter, the evidence that NGF and BDNF act as the pain mediators and modulators will be discussed (Fig. 1a). In addition, the clinical trials of management of chronic pain by neutralizing NGF are also introduced.



**Fig. 1** Summary of the mechanisms by which NGF exerts its neuromodulator in pain processing. (a) Schematic summary of pain processing from nerve terminals, the primary sensory neuron (DRG), and spinal cord (second-order neuron). (b) Acute phase of NGF exerting pain hypersensitivity through peripheral sensitization. In response to peripheral inflammation, NGF released from many types of cells sensitizes many membrane surface receptors (TRPV, ASIC, etc.) and releases pain mediators (SP, CGRP, and BDNF), resulting in mechanical allodynia and thermal hyperalgesia. (c) Later phase of NGF exerting pain hypersensitivity through peripheral and central sensitizations. Released NGF can transport to DRG through a retrograde transport way. In the DRG, many genes are activated and the increased pain mediators such as SP, BDNF, and TRPV can anterogradely transport to peripheral tissue or spinal cord which contributes to central sensitization. The process may take hours to days

# 2 Nerve Growth Factor

Nerve growth factor is the first identified neurotrophin and can be isolated into two forms, 7S or 2.5S (S is defined as the sedimentation coefficient) from male mouse salivary gland. Mature NGF is an approximately 12.5 kDa protein and usually forms homodimers after being cleaved from pro NGF which may have independent biological activity through activating p75<sup>NTR</sup>. NGF binds to TrkA, which directly

activates the downstream signaling (Levi-Montalcini 1998). The complexity of NGF-TrkA can be internalized and retrogradely transported to the cell bodies from the nerve terminals (Mantyh et al. 2011).

#### 2.1 Deprivation of NGF Results in Pain Insensitivity

NGF as a pain modulator has been identified by the finding that selective mutations in NGF or TrkA genes cause congenital insensitivity to pain in humans and loss of pain behaviors in genetically altered mice (Crowley et al. 1994; Einarsdottir et al. 2004). In NGF or TrkA knockout mice, most sensory neurons expressing TrkA die and are profoundly unresponsive to pain noxious stimuli (Crowley et al. 1994; Smeyne et al. 1994; Einarsdottir et al. 2004). In addition, patients with a mutation in TrkA gene have normal proprioception and normal sensation to innocuous pressure but present pain insensitivity to thermal stimuli (Indo 2002). Other studies observed that immunization against NGF in the critical period of the development, i.e., in the utero or prenatal period, results in depletion of substance P in dorsal root ganglia (DRG) neurons in animals (Ross et al. 1981; Johnson et al. 1989). Delivery of anti-NGF antibody in the early postnatal development causes a developmental switch of high-threshold mechanoreceptors to sensitive mechanoreceptors (Ritter et al. 1991).

Together, deprivation of NGF using genetic or immunological methods shows that NGF is essential for the survival of the sensory neurons in the prenatal development phase. However, NGF is not required for the survival of sensory neurons in the postnatal phase; instead, NGF mainly maintains the peptidergic phenotype of primary neuronal afferents.

# 2.2 NGF in Inflammatory Pain

Ample evidence shows that NGF is involved in the induction and persistence of various types of pain including inflammatory pain, cancer pain, neuropathic pain, and surgical pain. In these types of pathological pain, NGF and its high-affinity receptor TrkA are upregulated in the local region and DRG. Manipulating NGF or TrkA and downstream signaling could attenuate pain hypersensitivity in pathological pain. Given that the role of NGF signaling has been extensively studied in the inflammatory pain, thus the role of NGF as an inflammatory mediator that regulates pain process firstly and then its role in other types of pain will be discussed.

In inflammatory pain including carrageenan, formalin, and complete Freund's adjuvants as well as in models of autoimmune arthritis (Shelton et al. 2005) and ultraviolet B radiation-induced acute inflammation (Bishop et al. 2007), NGF and TrkA are increased in the inflamed tissue and DRG. The local increase of NGF is mainly released from the inflammatory cells, immune cells, Schwann cells, as well as noninflammatory cells such as keratinocytes and endothelial cells (Tron et al. 1990). The released NGF binds to TrkA on mast cells resulting in the release of inflammatory mediators such as histamine, serotonin (5-HT), and NGF

itself (Fig. 1b). When NGF binds to TrkA on the peptidergic fiber terminals, it activates the downstream intracellular signaling pathways, which subsequently activate or modulate many membrane surface receptors including bradykinin receptors; ion channels including transient receptor potential vanilloid 1 (TRPV1), acid-sensing ion channels (ASIC) 2/3, voltage-gated sodium (Na<sub>v</sub>) or calcium (Ca<sub>v</sub>) ion channels, and delayed rectified potassium ( $K^+$ ) currents; and putative mechanotransducers (Mantyh et al. 2011). The rapid modification of membrane surface receptors in the primary afferent terminals leads to sensitization of primary afferent terminals to mechanical or thermal stimuli (Bennett et al. 1998; Shu and Mendell 1999). The peripheral sensitization is rapid taking from minutes to hours (Fig. 1b). When the inflammation is persistent (e.g., inflammation induced by complete Freund's adjuvant). NGF and TrkA are transported to the DRG in a retrograde way (Ji et al. 2002). There NGF-TrkA signaling results in the production and release of neuropeptides such as substance P (SP) and calcitonin gene-related peptide (CGRP) and the other neurotrophin, BDNF, some of which are transported to peripheral tissue or the spinal cord (Lindsay and Harmar 1989; Woolf and Costigan 1999; Ji et al. 2002). The procedure of central sensitization may take several hours to days since it is involved in the retrograde transport, gene expression, anterograde transport, phenotype switch, and so on (Woolf and Costigan 1999) (Fig. 1c). A typical example is that NGF-induced pain hypersensitivity is selectively blocked by MK 801, a noncompetitive antagonist of N-methylp-aspartate (NMDA) receptor, which plays a critical role in the development of windup and central sensitization (Woolf 1996; Yaksh et al. 1999).

### 2.3 NGF in Other Types of Pain

NGF not only acts as an inflammatory mediator and results in peripheral and central sensitization in inflammatory pain but is also involved in other types of pain. In a mouse model of bone cancer, sustained delivery of anti-NGF antibody could inhibit the development of cancer pain, along with blocking the pathological sprouting of sensory and sympathetic nerve fibers, the formation of neuroma-like structures (Halvorson et al. 2005; Mantyh et al. 2010). This study suggests that NGF is released during the development of cancer pain. The released local NGF induces a remarkable reorganization of sensory and sympathetic nerve fibers, resulting in the formation of neuroma-like structure and sympathetic sprouting (Mantyh et al. 2010). Further study demonstrates that the source of NGF may be mainly from tumor-associated stromal, inflammatory, and immune cells (Ehrhard et al. 1993; Artico et al. 2008; Jimenez-Andrade et al. 2010). In this regard, injection of prostate cells which do not express detectable levels of NGF gene results in sclerotic bone lesions. In this model of bone cancer, TrkA-positive sensory and sympathetic nerve fibers are also observed to undergo the reorganization of pathological sprouting. Administration of anti-NGF to sequester the increased NGF in the early phase of prostate-induced bone cancer pain could block the nerve fibers sprouting and dramatically attenuate the development of cancer pain (Jimenez-Andrade et al. 2010). The findings that NGF contributes to the sprouting of sensory and sympathetic nerve fibers in cancer pain suggest that preemptive anti-NGF therapy not only attenuates the attendant pain but also inhibits the pathological changes of sensory and sympathetic nerve fibers, which may be critical for the development of chronic pain hypersensitivity. Therefore, anti-NGF therapy may be effective in treating various types of cancer pain which predicts the occurrence of tissue/nerve injury.

Recent studies have shown that NGF also regulates the postsurgical pain (Zahn et al. 2004; Wu et al. 2007). Hind paw incision (a common animal model of surgical pain) results in the increased NGF in the incised tissue, but not DRG. Anti-NGF treatment could attenuate the guarding pain- and heat hyperalgesia induced by hind paw incision although it would not affect the C-fiber sensitization (Zahn et al. 2004; Wu et al. 2007). Further study showed that the increased NGF after hind paw incision is mainly localized in nerve bundles including Schwann cells and axons (Wu et al. 2007). Thus, NGF may be taken up by the axon and transported into the DRG and exerts its biological function, similar to that in inflammatory pain. These findings suggest that NGF may be a potential target to treat postsurgical pain (Wu et al. 2007).

The role of NGF in the neuropathic pain has not been well established. Neuropathic pain is one of the greatest challenges in clinical practice and refractory to current therapies. Neuropathic pain is normally caused by nerve injury induced by various etiologies including diabetes, infection such as postherpetic neuralgia, drug treatment-induced neuropathy, and traumatic injury in the peripheral nerves or spinal cord (Pezet and McMahon 2006). In the traumatic injury-induced neuropathic pain models such as spinal nerve ligation or sciatic constriction injury, some axons have lesions while the other axons are intact. The role of NGF may be distinct in the development of this type of neuropathic pain. This is because in the injured axons, it undergoes Wallerian degeneration which may lead to the retrograde neurodegeneration to the spinal cord and loss of the contact with their peripheral target. NGF has been reported to exert neuroprotective effect on small-diameter nociceptive sensory neurons and therefore reverse sensory hyperalgesia (McMahon and Cafferty 2004). However, the clinical trial in diabetic neuropathy does not show any neuroprotective effect (Apfel 2002). In contrast, the injured Schwann cell or recruited immune cells may release NGF which would sensitize the intact axons or transport to the DRG in a retrograde way (Pezet and McMahon 2006). Thus, blocking NGF may be able to attenuate neuropathic pain. Several studies indeed have shown that anti-NGF could attenuate the pain hypersensitivity in rat models of sciatic constriction injury and spinal cord injury (Ramer and Bisby 1999; Deng et al. 2000; Gwak et al. 2003). Further studies are needed to elucidate how NGF is involved in the development of neuropathic pain.

#### 2.4 Anti-NGF Therapy in Clinical Trials

Ample experimental evidence has shown that blocking NGF or its high-affinity receptor TrkA could greatly inhibit the progress of chronic pain, in particular pain

induced by inflammatory injury. For example, systemic anti-NGF treatment prevents thermal and mechanical hyperalgesia in a complete Freund's adjuvantinduced inflammatory pain (Woolf et al. 1994). Administration of Trk-IgG fusion protein attenuates pain hypersensitivity induced by carrageenan (McMahon et al. 1995; Koltzenburg et al. 1999). In visceral inflammatory pain model, administration of anti-NGF antibody or Trk-IgG fusion protein markedly reduced the pain hypersensitivity. In addition, anti-NGF therapy also reversed the prolonged hyperalgesia in autoimmune arthritis (Shelton et al. 2005). Further studies have shown that anti-NGF therapy reduces pain caused by bone fracture or incisional pain suggesting that anti-NGF therapy is also effective for other types of pain (Jimenez-Andrade et al. 2007; Wu et al. 2007; Ghilardi et al. 2011).

The encouraging data of anti-NGF therapy in animal studies drive the scientists to investigate the role of NGF in humans. In the patient with chronic pain including arthritis, fibromyalgia, or peripheral nerve injury, NGF expression is increased locally (Aloe et al. 1992; Anand et al. 1997; Sarchielli et al. 2007). In addition, subcutaneous injection of NGF leads to persistent mechanical hyperalgesia in humans (Svensson et al. 2003; Rukwied et al. 2010). These studies suggest that increased local NGF may be involved in the development of chronic pain. There are several strategies to develop the anti-NGF-TrkA signaling for the treatment of chronic pain. These include NGF sequestration by anti-NGF antibody, inhibiting NGF binding to TrkA, blocking TrkA kinase activity by TrkA-Fc fusion protein (Shelton et al. 1995; Winston et al. 2003).

One typical molecule is a humanized anti-NGF monoclonal antibody tanezumab (RN-624, PF-04383119) (Cattaneo 2010). An early study has shown that, as compared to the placebo, tanezumab dose-dependently reduces the knee pain whilst walking in the patients with osteoarthritis, a form of arthritis in which the cartilage breaks down in the joint (Lane et al. 2010). The clinical trial of 450 patients shows that tanezumab along with NSAIDs reduces pain by 45–62 %, while a placebo only reduces pain by 22 % (Lane et al. 2010). In addition, tanezumab was also associated with dramatically greater improvement as measured by the patients' global assessment. However, phase 3 osteoarthritis trial was halted because 16 patients taking tanezumab were observed to have bone tissue death in their hip and knee joints and required the joint replacement (Garber 2011). Several possibilities are raised including the combined use of tanezumab and NSAIDs and reduced pain leading to much more movement of the joints (Lane et al. 2010; Garber 2011). Thus, it is still an important issue to address the safety of anti-NGF therapy in patients with chronic pain (Ackermann 2012).

#### 3 Brain-Derived Neurotrophic Factor

Another neurotrophin involved in pain is brain-derived neurotrophic factor (BDNF). BDNF is a 12.4 kDa basic protein and the second identified neurotrophin isolated from pig brain in 1982 (Barde et al. 1982). It is well known that BDNF plays a critical role for cell differentiation, neuronal survival, migration, dendritic

arborization, synaptogenesis, and activity-dependent synaptic plasticity in the central nervous system (Greenberg et al. 2009). Dysfunction of BDNF in the central nervous system would lead to cognitive dysfunction and anxiety/depression behaviors (Greenberg et al. 2009). There is ample evidence showing that BDNF also regulates pain processing as a central modulator (Pezet and McMahon 2006).

#### 3.1 Expression Pattern of BDNF and TrkB

Besides widely expressed in the brain, BDNF is also expressed in the tissues involving in the pain processing including the DRG and spinal cord. In the DRG, BDNF is constitutively expressed in the small- and medium-sized peptidergic neurons, which accounts for 10–40 % in the DRG neurons (Wetmore and Olson 1995; Luo et al. 2001). The synthesis of BDNF in the DRG neurons anterogradely transports into central terminals of primary afferent fibers in spinal cord dorsal horns. In the spinal cord, BDNF is mainly located in the large vesicles which also contain substance P, the common pain mediator. The BDNF contained in the sensory neurons and terminals in the spinal cord are believed to be released with activity. TrkB is also expressed in a subpopulation of DRG and the terminals in the dorsal horns of spinal cord (Zhou et al. 1993; Kashiba et al. 2003). In addition, TrkB is also expressed in nodose and trigeminal neurons and their terminals in trigeminal nucleus (Behnia et al. 2003). TrkB immunoreactive cells are also observed in the neurons of spinothalamic tract, which is associated with the transmission of painful information to supraspinal regions (Slack et al. 2005).

In addition to its expression in sensory neurons and central nervous system, BDNF gene is also expressed in the peripheral tissue. The expression of BDNF is higher in visceral tissues (such as the bladder, lung, and colon) than in the somatic tissues (Braun et al. 1999; Lommatzsch et al. 1999). In inflammatory or disease situations, BDNF is upregulated locally. For example, in the lung by allergen challenges, BDNF level is increased and expressed in the inflammatory infiltrates in subsets of T cells and the activated alveolar macrophages (Braun et al. 2004). Sequestration of the upregulated BDNF blocked the enhanced airway reactivity and persistent airway obstruction (Braun et al. 2004). In patients with chronic pancreatitis, BDNF is also greatly increased in ductural complexes and perineurium of the enlarged nerve. The increased level of BDNF is well correlated with the intensity of pain in the patients (Zhu et al. 2001).

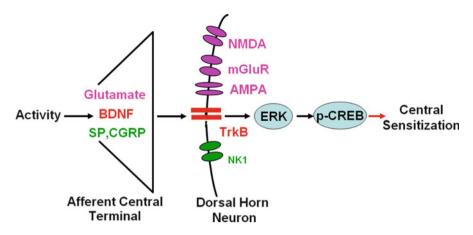
#### 3.2 Effect of Exogenous BDNF in Pain

There is accumulating evidence showing that BDNF regulates various types of pain states in different anatomical sites including peripheral tissues, DRG, spinal cord, and supraspinal levels. First, intrathecal delivery of exogenous BDNF could induce mechanical allodynia and hyperalgesia (Coull et al. 2005). Second, BDNF superfusion dramatically increases C-fiber-mediated reflexes which is greatly

inhibited in the BDNF-deficient mice (Kerr et al. 1999). In the inflammatory state, BDNF significantly increases the frequency of miniature excitatory postsynaptic currents (EPSCs) of the neurons in the spinal dorsal horns suggesting that BDNF activates nociceptive second-order neurons (Matayoshi et al. 2005). Third, BDNF-TrkB signaling activates the signaling closely associated with pain sensitization. For example, BDNF, after binding to TrkB, could potentiate glutaminergic neurotransmission, in particular N-methyl-D-aspartate (NMDA) receptor response (Slack et al. 2004). NMDA receptor is an ionotropic glutamate receptor and is believed to play a central role in central sensitization (South et al. 2003). Furthermore, intraspinal injection of BDNF can release the neuropeptide SP and induce the activation of extracellular regulated kinase (ERK) in the spinal cord (Kawasaki et al. 2004). Activation of ERK in the DRG and spinal cord has been reported to contribute to the development of pain such as inflammatory pain and neuropathic pain (Obata et al. 2004). BDNF may be able to activate cAMP response elementbinding protein (CREB) and early gene c-fos, both of which are sensitive and specific markers in the spinal cord for the noxious stimuli (Obata et al. 2004; Jongen et al. 2005). More recent study suggests that the increased BDNF in the spinal microglia induces the depolarizing shift in the anion reversal potential and subsequently converts the GABA ( $\gamma$ -aminobutyric acid) inhibitory neurons to excitatory neurons (Coull et al. 2005).

#### 3.3 Role of BDNF in Inflammatory Pain

As described above, pharmacological, biochemical, and neurophysiological evidence demonstrates that exogenous BDNF promotes pain processing. Extensive studies have shown that endogenous BDNF is increased in the DRG and spinal cord, and contributes to pain hypersensitivity in inflammatory pain. First, BDNF is increased in the DRG and spinal cord following peripheral inflammation including intraplantar formalin, CFA, and carrageenan (Pezet et al. 2002). The upregulation of BDNF is partly mediated by NGF since anti-NGF antibodies prevent the activation of BDNF in the spinal cord (Michael et al. 1997). In addition, a recent study also reports that TNF- $\alpha$  also contributes to the upregulation of BDNF and TrkB in the DRG (Onda et al. 2004). In DRG primary culture, chronic TNF-a treatment also upregulates the expression of BDNF and TrkB accompanied by the increased expression of CGRP and SP (Lin et al. 2011). In addition, tooth pulp inflammation increases BDNF expression in trigeminal ganglion (Kashiba et al. 2003). The increased BDNF is co-localized with CGRP and TRPV1 (Duric and McCarson 2007; Lin et al. 2011). Second, intrathecal injection of TrkB-IgG fusion protein dramatically ameliorates the pain behavior in response to peripheral inflammation suggesting that BDNF-TrkB signaling in the DRG and spinal cord contributes to inflammatory pain hypersensitivity (Lin et al. 2011). BDNF-TrkB signaling exerts their pro-nociceptive effect through multiple signaling such as activation of ERK1/2 signaling and CGRP, the well-known inflammatory pain mediator (Pezet and McMahon 2006) (Fig. 2).



**Fig. 2** Schematic summary of the mechanisms by which BDNF regulates pain processing as a central modulator in the spinal cord. Noxious stimuli induce the release of BDNF and neuro-transmitters including glutamate, SP, and CGRP. BDNF can bind to TrkB and subsequently activate ERK which contributes to the central sensitization. Glutamate can bind to its receptors including NMDA, mGluR, and AMPA receptors. BDNF also phosphorylates the subunit of NMDA receptor, NR1. SP binds its receptor, NK1. These pain neurotransmitters also play critical role for the central sensitization

BDNF-TrkB signaling is also observed in the spinal cord or peripheral tissue in the inflammatory visceral pain. In a bladder inflammation model, TrkB is increased in the bladder, pelvic ganglion, and DRG (Qiao and Vizzard 2002). In addition, BDNF level is increased in the rectosigmoid biopsies which correlates with the pain score in the patients with irritable bowel syndrome (Yu et al. 2012). In experimental inflammation of mouse gut, BDNF is upregulated in the DRG in the experimental colitis (Yang et al. 2010). The pain intensity in this inflammatory visceral pain is reduced in BDNF (+/-) mice as compared with BDNF (+/+) mice (Yu et al. 2012). Our recent study showed that the modulation of BDNF in visceral pain is sex specific. Anti-BDNF antibody pretreatment attenuates the nocifensive response in the males, whilst exacerbates the nocifensive response in the females in visceral pain induced by acetic acid intraperitoneal injection (Li et al. 2010). The upregulation of BDNF in the pancreas has also been reported in the patients with chronic pancreatitis, and the level of BDNF correlated with pain intensity (Zhu et al. 2001). All of these studies strongly indicate that BDNF regulates inflammatory pain through peripheral and central mechanisms.

#### 3.4 Role of BDNF in Neuropathic Pain and Surgical Pain

In neuropathic pain, BDNF expression is also dysregulated in the sensory neurons. However, the expression of BDNF in the sensory neurons is different in the different neuropathic pain models. For example, in the axotomy model induced by sciatic nerve section, BDNF level is upregulated in the medium- and large-diameter DRG

neurons and the central terminals in the deep layers of the dorsal horns in the spinal cord (Michael et al. 1999). However, the expression of BDNF in the small-diameter neurons is decreased probably due to the neurodegeneration induced by the axotomy. In another neuropathic pain model induced by sciatic nerve constriction injury, BDNF is upregulated in the small-diameter neurons. The increased BDNF in the small-diameter neurons may be mediated by  $TNF-\alpha$  through inducing NGF expression (Onda et al. 2004). A series of studies have shown that the activated microglia can release BDNF in the spinal nerve ligation-induced neuropathic pain model (Onda et al. 2004). The increased BDNF in microglia activates TrkB, which, subsequently, converts inhibitory neurons to excitatory function through inverts the polarity of currents induced by  $\gamma$ -aminobutyric acid (GABA) (Coull et al. 2005). The disinhibition of GABA receptor-mediated neurons would contribute to the peripheral nerve injury-induced tactile allodynia (Coull et al. 2005). The upstream signaling of microglial BDNF includes ATP receptors, in particular P2X4 receptors through P2X4R-evoked increase in Ca<sup>(2+)</sup> and the activation of p38 mitogen activation protein kinase (Trang et al. 2009a). Thus, in peripheral nerve injury-induced neuropathic pain, the upregulation BDNF can be from the neurons or microglia and maintains the persistent pain hypersensitivity after nerve injury.

Our previous study has shown that BDNF is also increased in the DRG and spinal cord after surgical incision. The increased BDNF is mainly localized in the neurons as well as glial cells. Intrathecal delivery of anti-BDNF could dramatically reduce the incision-evoked pain hypersensitivity suggesting that BDNF in the DRG and spinal cord also contribute to pain hypersensitivity in surgical pain (Li et al. 2008). BDNF is also reported to be involved in cancer-induced pain through modulating NMDA receptors (Wang et al. 2012). Minocycline, a potent microglia inhibitor, could prevent the activation of BDNF in the model of bone cancer pain (Wang et al. 2012). This suggests that BDNF released from microglia contributes to cancer-induced bone pain.

#### 3.5 Antinociceptive Effect of BDNF

Interestingly, in inflammatory pain, the expression of BDNF in the hippocampus is downregulated (Duric and McCarson 2007). Midbrain injection of BDNF, but not NGF, significantly exerts analgesia by elevating the tail-flick response latency and the latency in the hot-plate experiment (Siuciak et al. 1994, 1995). In addition, midbrain injection of BDNF also reduces the paw flinch response in the formalin intraplantar injection-induced pain (Siuciak et al. 1994, 1995). The analgesic effect of BDNF in the midbrain is associated with serotonergic neurons and can be prevented by naloxone (Siuciak et al. 1994, 1995). In addition, heterozygous BDNF(+/–) mice develops the hyperalgesia accompanied with the premature age-associated decreased forebrain 5-HT level and fiber density (Lyons et al. 1999). These findings suggest that supraspinal BDNF may exert antinociceptive effect through regulating the function of serotonergic neurons and levels of endogenous opioid receptors.

# 4 The Involvement of P75<sup>NTR</sup> in Pain

Unlike TrkA and TrkB,  $p75^{\text{NTR}}$  is a low-affinity binding receptor for the whole family of neurotrophins. However,  $p75^{\text{NTR}}$  is the high-affinity binding receptor of precursor of neurotrophins such as pro NGF or proBDNF (precursor of BDNF).  $p75^{\text{NTR}}$  has been reported to exert multiple neuronal functions in the presence of Trk receptors (see review (Lu et al. 2005)). In the absence of Trks, neurotrophin acts on  $p75^{\text{NTR}}$  and promotes apoptosis. In a neuropathic pain model,  $p75^{\text{NTR}}$  expression is increased in the axon transport and uninjured primary afferents (Obata et al. 2006). Functional inhibition of  $p75^{\text{NTR}}$  in the DRG could reverse neuropathic pain after nerve injury. Further study shows that  $p75^{\text{NTR}}$  promotes neuropathic pain through enhancing TrkA signaling and activating p38 mitogenactivated protein kinase and TPRV1. These findings indicate that  $p75^{\text{NTR}}$  contributes to nerve injury-induced neuropathic pain (Obata et al. 2006). More recent study shows that morphine tolerance is attenuated in the wild-type but not  $p75^{\text{NTR}}$  (-/-) mice in which the induction of opioid physical dependence is not developed.

### 5 Conclusion

Neurotrophins, in particular NGF and BDNF, regulate and drive acute and chronic pain as pain modulators in adults. NGF contributes to pain processing mainly acting as peripheral inflammatory mediators. It can directly sensitize the peripheral nociceptive afferents and lead to peripheral sensitization rapidly. It can also retrogradely transport to sensory neurons and activate downstream signaling and gene expression. Anti-NGF therapy for treatment of chronic pain represents a novel class of analgesic therapy, but its safety is still a major challenge for the clinical application.

BDNF regulates pain processing as central modulators. BDNF is increased in the primary sensory neurons and spinal cord in the various types of pain. BDNF binds to TrkB and activates many pain mediators such as NMDA receptors, neuropeptides, and ERK, leading to the central sensitization. Sequestration of the increased BDNF through intrathecal delivery attenuates most types of pain. BDNF in the supraspinal level is downregulated in inflammatory pain, and midbrain infusion of BDNF exerts analgesic effect.

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# **Neurotrophin Signaling in Cancer**

Fei Tan, Carol J. Thiele, and Zhijie Li

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#### Abstract

Neurotrophins are growth factors that play a major role in neuron survival, proliferation, differentiation, and apoptosis. There are four neurotrophins, nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), neurotrophin-3 (NT-3), and neurotrophin-4/5 (NT-4/5) which function by interacting with Trk (tropomyosin-related kinase) tyrosine kinase receptors,

F. Tan (🖂)

C.J. Thiele • Z. Li

Department of Neurology, Shengjing Hospital of China Medical University, Shenyang, China e-mail: tanf@sj-hospital.org

Cellular and Molecular Biology, Pediatric Oncology Branch, National Cancer Institute, National Institute of Health, Bethesda, MD, USA e-mail: ct47a@nih.gov; lizhijie68@hotmail.com

TrkA, TrkB, TrkC, or p75NTR receptor. Binding of neurotrophins with their cognate Trk receptors activates Ras/mitogen-activated protein kinase (MAPK) pathway, Phosphatidylinositol-3 kinase (PI3K)/Akt pathway, and Phospholipase C- $\gamma$  (PLC- $\gamma$ ) pathway which are involved in cell survival, proliferation, differentiation, and apoptosis. Activation of the pan-neurotrophin receptor, p75NTR, leads to JNK pathway activation that induces cell apoptosis or activates a NF-KB cell survival pathway. Although neurotrophins were originally found to act on neurons, studies indicate they also have activities in nonneuronal cells. Trk activation or mutation occurs in tumors of neuronal origin, like neuroblastoma and medulloblastoma, as well as in nonneuronal tumors like thyroid, breast, lung, and prostate cancer. In neuroblastoma, expression of TrkA occurs in tumors that are biologically favorable and prone to spontaneous regression or differentiation which may be due to the absence or presence of its ligand (NGF) in the microenvironment. Expression of TrkB and/or its ligand BDNF in neuroblastoma tumors is often associated with chemo-resistance, metastasis, and a poor prognosis. In breast cancer, overexpression of TrkA promotes cell growth, migration and invasion in vitro or tumor growth, metastasis, and angiogenesis. Based on these studies, small molecule inhibitors targeting Trk receptors or downstream targets in their signaling pathways have been developed. The Trk selective inhibitor CEP-701 (Cephalon) or the Akt inhibitor perifosine are both in phase 1 and 2 clinical trials. The effects of these small molecule inhibitors, alone or in combination with chemotherapeutic drugs, are discussed.

List of Abbrevia	ations
BDNF	Brain-derived neurotrophic factor
ERK	Extracellular signal-regulated kinase
Grb2	Growth factor receptor-bound protein 2
HGF	Hepatocyte growth factor
JNK	c-Jun N-terminal kinase
MAPK	Mitogen-activated protein kinases
MEK	MAP kinase kinase
mTOR	Mammalian target of rapamycin
NGF	Nerve growth factor
NSCLC	Non-small cell lung cancer
NT-3	Neurotrophin 3
NT-4/5	Neurotrophin 4/5
P75NTR	P75 neurotrophin receptor
PI3K	Phosphatidylinositol-3 kinase
PLC-γ	Phospholipase C-γ
PTB domain	Phosphotyrosine-binding domain
SCLC	Small cell lung cancer
Shc	Src-homology collagen protein
SOS	Son of sevenless
Trk	Tropomyosin-related kinase

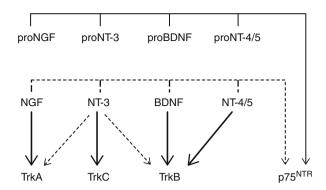
#### 1 Introduction

Neurotrophin are growth factors implicated in several different functions in the nervous system, including survival, proliferation, differentiation, myelination, apoptosis, axonal growth, and synaptic plasticity. In mammals, the neurotrophin family consists of nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), neurotrophin 3 (NT-3), and neurotrophin 4/5 (NT-4/5) (Snider 1994). With the exception of NT-4/5, which is not detected in avian species, the neurotrophin sequences are evolutionarily conserved in vertebrates. The *nt-6* and *nt-7* genes encode neurotrophins NT-6 and NT-7 and have been identified only in fish, and do not appear to have mammalian orthologs (Gotz et al. 1994; Nilsson et al. 1998).

The signal transduction pathways that mediate the diverse biological functions of the neurotrophins are initiated through interactions with two categories of cell surface receptors: the selective tropomyosin-related kinase (Trk) tyrosine kinase receptors and the "shared" p75 neurotrophin receptor (p75NTR). These two types of receptors share no sequence similarity in either ligand-binding or cytoplasmic domains, and activate distinct neurotrophin-dependent signaling pathways. In many instances, Trk receptors and p75NTR not only activate autonomous pathways but also collaborate to mediate effects of the neurotrophins. There are three Trk receptors: TrkA, TrkB, TrkC. The Trk receptor was originally discovered as a rearrangement of non-muscle tropomyosin and a then unknown tyrosine kinase. This tyrosine kinase was referred to as Trk and subsequently identified as a receptor for NGF. Trk became TrkA with the indentification of TrkB and TrkC which were based on their DNA sequence similarity to Trk (Hempstead et al. 1991; Klein et al. 1989, 1991; Lamballe et al. 1991). Each neurotrophin binds with high affinity to Trk receptors (Fig. 1): NGF to TrkA, BDNF to TrkB, NT-3 to TrkC. NT-4/5 binds TrkB, too. NT-3 also binds to TrkA and TrkB receptors that contain an additional short insert in the extracellular domain (Clary and Reichardt 1994; Strohmaier et al. 1996). The p75NTR belongs to the tumor necrosis factor (TNF) superfamily and was the first identified receptor for NGF (Chao et al. 1986; Radeke et al. 1987). All neurotrophins bind to the common p75NTR receptor with low affinity. Recently, pro-neurotrophins (the precursors of the neurotrophins: pro-NGF, pro-BDNF, pro-NT-3, pro-NT-4/5) have been identified as high-affinity ligands for p75NTR in complex with a co-receptor, sortilin (Lee et al. 2001; Nykjaer et al. 2004; Teng et al. 2005) (Fig. 1).

Disturbances in neurotrophins, their receptors, or neurotrophin/receptor-initiated signaling pathways have been associated not only with various neurodegenerative disorders (e.g., Alzheimer's disease, Parkinson's disease), but also to several types cancers of neuronal (e.g., neuroblastoma, medulloblastoma, gliomas) and nonneuronal origin (e.g., breast, lung, prostate cancers).

This chapter will describe neurotrophins, their receptors, and the signaling pathways initiated by binding of neurotrophins to Trk or p75NTR receptors. The expression of neurotrophins and/or their receptors in neuronal and nonneuronal cancers, as well as the potential to target these neurotrophins/receptors signaling pathways in the treatment of cancers, will be described.



**Fig. 1** Neurotrophins and their receptors: Neurotrophins preferentially bind to Trk receptors: NGF binds specifically to TrkA; BDNF and NT-4/5 recognize TrkB; NT-3 activates TrkC. NT-3 is also able to activate TrkA and TrkB with less efficiency. Neurotrophins bind p75NTR with equal affinity. Pro-neurotrophins bind p75NTR with high affinity

#### 2 Neurotrophins

NGF was the first member of the neurotrophin family to be identified. NGF activity was initially observed in a mouse sarcoma that induced the dramatic growth of peripheral nerve fibers when transplanted into chick embryos. NGF was purified from mouse submandibular gland, which expressed extremely high levels of the factor. A dimer of 13.3 kDa polypeptides, termed the  $\beta$  subunit, was found to contain the biological activity of NGF. The  $\beta$ -NGF polypeptide chain is produced as a larger propeptide that is proteolytically cleaved to generate the mature form of a homodimeric polypeptide of about 26 kDa. A major biological function of NGF is the maintenance and survival of post-mitotic neurons; it is normally produced by the synaptic target field in limiting amounts to attract susceptible growth cones to functional site. NGF is a major player in the development of several peripheral nervous system subpopulations, as well as several central nervous system regions; as such, it has been considered a strong candidate for the treatment of several neurodegenerative conditions. In addition to its role in the maturation and maintenance or neuronal cells, NGF also has significant effects on nonneuronal cells. For example, it triggers chemotaxis in melanocytes, inhibits the migration of Schwann cells, has been reported to stimulate the proliferation of chromaffin cells, and is also an autocrine survival factor for B lymphocytes (Butte 2001; Reichardt 2006; Sofroniew et al. 2001; Wiesmann and de Vos 2001).

BDNF was the second neurotrophin to be discovered after its purification from pig brain in 1982. BDNF was produced initially as pro-BDNF (M.W.  $\sim$ 30 kDa), then cleaved to generate the mature form (M.W.  $\sim$ 14 kDa) which shared about 50 % amino acid identity with NGF. BDNF has survival- and growth-promoting actions on a variety of neurons, including dorsal root ganglion cells and hippocampal and cortical neurons. Certain peripheral sensory neurons, especially those in vestibular and nodose-petrosal ganglia, depend on

the presence of BDNF because BDNF homozygous (-/-) knockout mice lack these neurons. Unlike NGF, neither sympathetic or motor neurons are affected by BDNF. BDNF also plays a role in brain development and synaptic plasticity (Binder and Scharfman 2004; Reichardt 2006). As neural crest cells coalesce to form sympathetic ganglia, TrkB-positive cells are seen in both chicken and mouse embryos. The coexpression of TrkB and HNK-1 confirms that they are migrating neural crest cells. Subsequently, TrkB-positive cells begin to express the neural-specific markers Hu C/D and Islet-1, eventually commence neural differentiation (Straub et al. 2007).

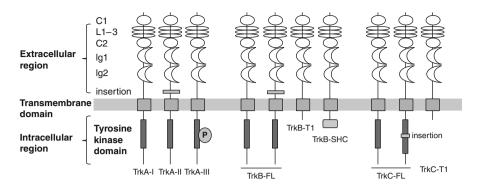
NT-3 was identified using degenerate oligonucleotides to amplify by PCR novel cDNA sequences related to NGF and BDNF. The NT-3 cDNA encoded a mature polypeptide of predicted molecular weight 13.6 kDa. Recombinant NT-3 supports the survival of and neurite outgrowth from chick embryonic neurons, a result that established its neurotrophic activity. A PCR strategy similar to that used for NT-3 was used to identify a fourth mammalian neurotrophin, NT-4 (also known as NT-4/5), which was also shown to possess neurotrophic activity (Reichardt 2006).

## **3** Neurotrophin Receptors

#### 3.1 Trk Receptors

Trk receptors are prototypical receptor tyrosine kinases that contain an extracellular domain, a transmembrane domain, and an intracellular domain. The extracellular domain is composed of a cysteine-rich domain followed by three leucine-rich domain, another cysteine-rich domain, and two immunoglobulin-like C2 type domains (Ig-C2). The cytoplasmic region contains a kinase domain (Fig. 2)(Brodeur et al. 2009; Gills and Dennis 2009; Reichardt 2006; Thiele et al. 2009). Binding of neurotrophins to Trk receptors occurs mainly through the Ig-C2 domains, with the domain closer to the transmembrane region playing a more prominent role in binding. The leucine-rich domains and the cysteine domains may also be involved in binding with neurotrophins. In addition to ligand binding, the Ig-C2 domains can also stabilize the monomeric form of the Trk receptor to prevent spontaneous dimerization and activation in the absence of neurotrophins. Deletions, chimeric receptors, or point mutations that disrupt the structure of the first or second Ig-C2 domain allow ligand-independent receptor activation. Differential splicing that leads to a loss of the first Ig-C2 domain of TrkA exhibits ligand-independent signaling and transformation activity when transfected into NIH-3T3 cells (Tacconelli et al. 2004).

The generation of alternatively spliced isoforms of Trk genes leads to functionally distinct receptors (Fig. 2). TrkA has three isoforms, TrkA-I, TrkA-II, and TrkA-III. TrkA-II is the isoform which is expressed in neuronal cells. TrkA-I is a nonneuronal TrkA isoform that lacks exon 9. Exon 9 encodes 6 amino acids in the extracellular domain near the transmembrane region (insertion) in TrkA and does not alter the reading frame. This does not affect NGF binding or receptor activation, but the TrkA-II isoform has enhanced responsiveness to NT3. TrkA



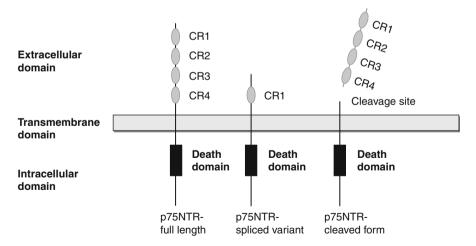
**Fig. 2 Structures of the Trk receptors**: The domain structures of the Trk receptors are schematized. Each Trk receptor has an extracellular domain, transmembrane domain, and an intracellular tyrosine kinase domain. The locations of two cysteine-rich domains (C1, C2), three leucine-rich domains (L1-3), and two immunoglobulin-like C2 type domains (Ig1, Ig2) in the extracellular domain of Trk receptors are depicted. There is an insert in the extracellular domain near the transmembrane region in TrkA or TrkB receptor. The insert of TrkC receptor is in the intracellular tyrosine kinase domain. Several truncated isoforms of TrkB and TrkC that lack the tyrosine kinase domain have been identified

isoforms lacking this sequence interact normally with NGF but show reduced NT-3 binding (Clary and Reichardt 1994). There is also an isoform, TrkA-III that is expressed early during development and splices out exons 6, 7, and 9. TrkA-III is constitutively active, independent of ligand (Tacconelli et al. 2004). Enhanced expression of TrkAIII occurs under hypoxia. TrkB is expressed as both full-length (kinase-intact) and truncated (kinase-deleted) isoforms. Full-length TrkB receptors have alternatively spliced product lacking 6–9 amino acids in the extracellular juxtamembrane region, like TrkA. TrkB variants lacking this region bind BDNF normally but show decreased affinity for NT-3 and NT-4/5. There are at least two kinase-deleted isoforms of TrkB that are truncated just beyond the transmembrane domain and differ by the absence (TrkB-T1) or presence (TrkB-SHC) of the SHC-binding site (Brodeur et al. 2009; Reichardt 2006; Thiele et al. 2009). Studies indicating that expression of truncated TrkB receptors inhibits BDNF induced neurite activity suggest that such isoforms may act as dominantnegative inhibitors of the TrkB kinase. This inhibition may occur by ligand sequestration or by formation of functionally inactive heterodimers. TrkC has full-length and truncated isoforms. Full-length TrkC receptors can be produced with amino acid inserts (14, 25, or 39 residues) in their kinase domain. Forms of TrkC containing insertions within the kinase domain retain their ability to autophosphorylate in response to neurotrophin-3, but cannot mediate proliferation in fibroblasts or neuronal differentiation in PC12 cells (Valenzuela et al. 1993). Truncated TrkC is the isoform without the intracellular kinase domain which may act as a dominant-negative inhibitor of kinase-intact TrkC. Thus, alternative splicing of the Trk receptors affects the specificity of neuronal responsiveness to neurotrophins.

## 3.2 p75NTR receptor

p75NTR is a 399-amino acid Type I transmembrane receptor which contains an extracellular, transmembrane, and intracellular domain. The extracellular domain contains four cysteine-rich domains (CR1-4). The intracellular domain contains a death domain which made it the first identified but last accepted member into the Death Receptor Family (TNF and TRAIL) (Fig. 3). Experimental and structural modeling studies have suggested that CR3 may be primarily responsible for interaction with neurotrophins (Yan and Chao 1991). p75NTR has both full-length and truncated isoforms. The p75ntr gene gives rise to a full-length receptor (p75NTR. FL) and an alternatively spliced isoform (p75NTR.T) lacking exon 3 (CR2-4). The full-length p75NTR receptor is also cleaved by a constitutively active metalloproteinase to generate a soluble extracellular domain that is capable of binding neurotrophins, and a receptor fragment containing transmembrane and intracellular domain (Fig. 3) (Zupan et al. 1989). The soluble form of p75NTR is produced at very high levels during development and following peripheral nerve injury, but the biological role(s) of the proteolytic products remain poorly defined (Zupan et al. 1989).

All neurotrophins bind to p75NTR receptor with similar affinity. The affinity of neurotrophins to TrkA and TrkB receptors is modulated by the presence of p75NTR, which enhances specificity for their primary ligands NGF and BDNF, respectively (Benedetti et al. 1993). Formation of high-affinity binding sites for NGF requires the presence of a correct ratio of full p75NTR and Trk receptors, as mutations of the cytoplasmic or transmembrane domains of either TrkA or p75NTR abolish the configuration of high-affinity binding sites. At the same time, p75NTR reduces the ability of NT-3 to activate TrkA, and NT-3 and NT-4 to activate TrkB.



**Fig. 3 Structure of p75NTR**: p75NTR has an extracellular domain which consists of four cysteine-rich domain (CR1, CR2, CR3, CR4), a transmembrane domain, and an intracellular death domain. p75NTR has a spliced variant that lost the CR2 to CR4, and a cleaved form

This effect has been recently demonstrated in vivo in sympathetic neurons, where NT-3 can signal locally via TrkA receptors in the absence of p75NTR. Once axons reach their final target and receive NGF, TrkA retrograde signaling upregulates p75NTR, and the cells no longer respond to NT-3.

# 4 Neurotrophin Signaling Pathways

# 4.1 Neurotrophin Signaling Pathways Mediated by Trk Receptors

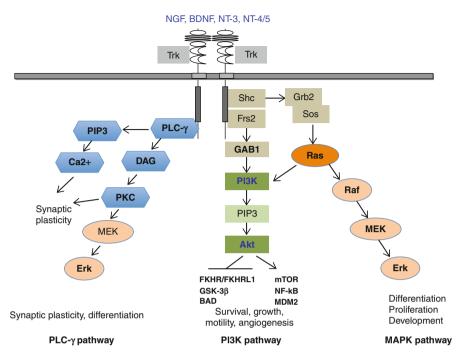
Binding of neurotrophin homodimers causes receptor dimerization. autophosphorylation on tyrosine (Y) residues within the activation loop (Y670, Y674, and Y675 relative to human TrkA), recruitment of different cytosolic adaptor proteins and enzymes which leads to the activation of several different signaling pathways. Among the signaling pathways activated by Trk receptors in response to neurotrophins, the Ras/mitogen-activated protein kinase (MAPK) pathway, phosphatidylinositol-3 kinase (PI3K) pathway, and PLCy-protein kinase C (PLC $\gamma$ ) pathways are primarily studied (Fig. 4) (Kaplan and Miller 2000; Reichardt 2006). Most of the work describing these signaling pathways has been performed in PC12 cells, a rat adrenal pheochromocytoma cell line that responds to NGF and in many respects resembles sympathetic neurons. These signaling pathways have also been studied largely in the context of NGF activation of TrkA and extrapolated to the other Trk receptors.

#### 4.1.1 Ras/MAPK Signaling Pathway

After neurotrophins, Trk receptor tyrosine binding to kinases are autophosphorylated and activated which recruits the adaptor protein Shc (Srchomology collagen protein) to phospho-Y490 on Trk receptors via interactions with the Shc PTB (phosphotyrosine binding) domain. Trk-mediated phosphorylation of Shc creates a phosphotyrosine site on Shc that recruits the adaptor Grb2 (growth factor receptor-bound protein 2), which is bound to the Ras exchange factor son of sevenless (SOS). These adaptors promote Ras activation which phosphorylates and activates Raf. Raf then phosphorylates and activates MEK (MAP kinase kinase), which in turn induces the phosphorylation and activation of ERK (extracellular signal-regulated kinase) (Fig. 4). Activated ERK translocates to the nucleus and transactivates transcription factors such as STAT1/3, Elk1, Myc, modulating gene expression to promote growth, differentiation, or development (Arevalo and Wu 2006; Reichardt 2006).

#### 4.1.2 PI3K/Akt Pathway

PI3-kinases are activated through Ras-dependent and Ras-independent pathways, and the comparative importance of these two mechanisms differs in various subsets of neurons. In many neurons, Ras-mediated activation of PI3-kinase initiates the major pathways through which survival signals are conveyed. In addition, PI3-kinase can



**Fig. 4** Neurotrophin signaling pathways mediated by Trk receptors: Binding of neurotrophins to their Trk receptors leads to dimerization of the receptors, activation of tyrosine kinases, and the recruitment of different adaptors proteins and enzymes, resulting in activation of several signaling pathways, including Ras/MAPK pathway, PI3K/Akt pathway, and PLC-γ pathway. These signaling pathways are involved in the cell differentiation, proliferation, survival, motility, and synaptic plasticity

also be activated through Ras-independent pathways. Recruitment of Gab1 and Frs2 permits subsequent binding and activation of PI3K. Activated PI3-kinase generates phosphatidylinositol 3.4-bisphosphate (PIP2). PIP2 acts as a substrate to generate the second messenger phosphatidylinositol-3,4,5-triphosphate (PIP3), which recruits and binds AKT to the cell membrane via its pleckstrin homology domain. Akt is then phosphorylated and activated by PDK1 and PDK2. Phosphorylated Akt is translocated to the cytoplasm, nucleus, or mitochondria where it functions to phosphorylate downstream targets, leading to their activation or inactivation. Akt inactivates the FKHR/FKHRL1 (FOXO1) transcription factor, glycogen synthesis kinase  $(GSK-3\beta)$ , and the Bcl-2 pro-apoptotic family member BAD known to play a proapoptotic role. Akt activates a number of proteins important for cell survival such as mTOR (mammalian target of rapamycin) which is important in protein synthesis, MDM2 which is an important negative regulator of the p53 tumor suppressor, and the transcription factor NF- $\kappa$ B (Fig. 4) (Arevalo and Wu 2006; Brodeur et al. 2009; Reichardt 2006; Thiele et al. 2009). Aside from a role in cell growth and survival, neurotrophin-mediated activation of the PI3K signaling pathway stimulates cell motility and increases production of VEGF. BDNF-mediated activation of TrkB in cerebellar granule neurons, as well as NGF-mediated activation of TrkA in dorsal root ganglion neurons, stimulates an increased production of VEGF that are mediated by increases in HIF- $\alpha$  (Nakamura et al. 2011). Neurotrophin-stimulated increases in VEGF may mediate communication between the vasculature and neurons during development, cell stress, or hypoxia.

#### 4.1.3 PLC-γ Pathway

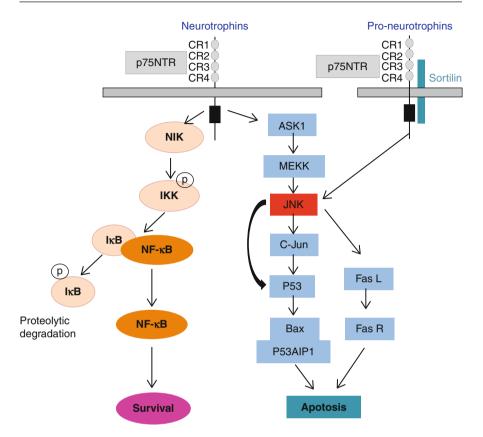
Following Trk kinase activation, PLC- $\gamma$  is recruited to a docking site surrounding phosphorylated Y785 on TrkA and to similar sites on TrkB and TrkC. Docked PLC- $\gamma$  is activated through Trk-mediated phosphorylation and then hydrolyzes PIP2 to generate PIP3 and DAG. The presence of PIP3 results in Ca2+ release from cytoplasmic stores. DAG stimulates DAG-regulated isoforms of protein kinase C. Together, these signaling molecules activate a number of intracellular enzymes, including almost all PKC isoforms, Ca2+ -calmodulin-dependent protein kinases, and other Ca2+ -calmodulin-regulated targets. The proteins activated as a result of PLC- $\gamma$  activation include PKC- $\delta$ , which is required for NGF-induced activation of MEK1 and Erk (Fig. 4). Signaling through this pathway controls expression and/or activity of many proteins, including ion channels and transcription factors (Arevalo and Wu 2006; Brodeur et al. 2009; Reichardt 2006; Thiele et al. 2009).

# 4.2 Neurotrophin Signaling Pathways Mediated by p75NTR Receptor

Binding of p75NTR to neurotrophins initiates opposing pathways, the pro-apoptotic signaling pathway and the pro-survival signaling pathway. The pro-survival effect of p75NTR/neurotrophins is mediated by the activation of the transcription factor, NF- $\kappa$ B (Karin and Lin 2002). The prototypic NF- $\kappa$ B molecule consists of a p65/p50 heterodimer, which is normally sequestered in the cytoplasm via binding to the inhibitory protein I $\kappa$ B. In response to neurotrophin binding to p75NTR, I $\kappa$ B is phosphorylated by the upstream IKK complex, leading to ubiquitination and degradation. Without the inhibition of I $\kappa$ B, NF- $\kappa$ B translocates into the nucleus where it regulates transcription of target genes (Fig. 5). Among the genes induced by NF- $\kappa$ B are several that promote survival, including members of the inhibitor of apoptosis family, IAP, and Bcl-2 homologs (Pahl 1999).

The pro-apoptotic effecter of activated neurotrophin/p75NTR is the JNK (c-Jun N-terminal kinase) signaling pathway. Activation of p75NTR induces activation of ASK1 and MEKK which in turn activates JNK. JNK either directly, or through the phosphorylation of c-jun, activates p53 which induces apoptosis by regulating its targets such as pro-apoptotic Bcl-2 family members Bax, Bad, Puma. In addition, activation of JNK induces expression of Fas ligand in neuronal cells, promoting apoptosis through activation of the Fas receptor (Fig. 5).

As mentioned early, p75NTR binds the unprocessed pro-neurotrophins with high affinity, stimulating an apoptotic signal. The pro-neurotrophins actually bind with high affinity to a complex of p75NTR with sortilin, a Vps10-domain



**Fig. 5** Neurotrophin signaling pathways mediated by p75NTR receptor: Binding of neurotrophins to p75NTR either activates NF- $\kappa$ B signaling pathway to mediate cell survival, or activates JNK signaling pathway to induce cell apoptosis. Pro-neurotrophins bind to p75NTR to initiate cell apoptosis via the JNK signaling pathway

containing protein. Both sortilin and p75NTR participate directly in binding to the pro-neurotrophins. The presence of sortilin is required to generate apoptosis following engagement of p75NTR by the pro-neurotrophins. The JNK signaling pathway mediates the p75NTR/pro-neurotrophins-induced apoptosis.

# 5 Neurotrophins and Their Receptors in Cancers

## 5.1 Neurotrophins and Their Receptors in Neuronal Origin Cancers

**Neuroblastoma** is the most common extracranial solid tumor in children. It arises from neural crest precursor cells. The correlation between the expression levels of neurotrophins/neurotrophin receptors in tumors and the biological/clinical

characteristics of tumors has been extensively studied in neuroblastoma. The prognosis of neuroblastoma patients is varied. In some neuroblastoma patients, the tumors can regress or are very sensitive to the current cytotoxic regimens and they have a 90 % overall survival. In contrast, some neuroblastoma patients have tumors that are very resistant to chemotherapy, readily metastasize, and these patients have a poor prognosis. Differential Trk/neurotrophin expression has been used to correlate these clinical features.

TrkA (I, II) is expressed at high levels in biologically favorable neuroblastoma, and associated with favorable clinical features, so TrkA expression in tumors of neuroblastoma patients becomes a strong predictor of good prognosis (Nakagawara et al. 1993; Suzuki et al. 1993). Consistent with this observation, high TrkA expression in neuroblastoma cell lines is associated with decreased invasiveness, slow tumor cell growth, and low expression of angiogenic growth factors (Eggert et al. 2002; Lucarelli et al. 1997). Low-stage neuroblastomas usually express high TrkA and respond to NGF by enhanced survival and terminal differentiation (Nakagawara et al. 1993). TrkA expression is low or absent in most advanced stage tumors, and they do not undergo terminal differentiation in response to NGF, consistent with a model in which the NGF/TrkA pathway is responsible for differentiation and/or regression of favorable neuroblastoma. However, the recently identified isoform of TrkA (TrkA-III) was found to be more highly expressed in neuroblastoma tumors of patients with a poor prognosis (Tacconelli et al. 2004).

Most neuroblastomas express very low or undetectable levels of NGF. A limited amount of NGF may be supplied from the stromal-infiltrating cells such as schwannian cells and fibroblasts. One model posits that in regressing neuroblastoma, tumor cells expressing TrkA receptor are dependent on limiting amounts of NGF which would lead to differentiation in the presence of NGF or apoptosis in its absence. This model is based on the observation that normal developing sympathetic neurons survive and differentiate in the presence of target-derived supplement of neurotrophins or die in its absence (Nakagawara 1998a, b). Recently, a study found CCM2 as a key mediator of TrkA-dependent cell death. CCM2 is the protein product of the cerebral cavernous malformation 2 gene, it interacts with the juxtamembrane region of TrkA via its PTB (phosphotyrosine binding) domain. Coexpression with TrkA, CCM2 mediates TrkA-induced cell death, while CCM2 knockdown attenuates TrkA-dependent cell death. Furthermore, increased TrkA and CCM2 transcript expression correlates strikingly with positive clinical outcome in a large cohort of neuroblastoma patients (Harel et al. 2009).

Full-length TrkB and BDNF are expressed individually or co-expressed in more aggressive, unfavorable neuroblastoma (Nakagawara et al. 1994), Whereas truncated TrkB is expressed in more differentiated tumors, such as ganglioneuroblastoma. Addition of BDNF to the TrkB-expressing neuroblastoma cells enhanced the survival of cells exposed to serum-free media (Nakagawara et al. 1994), or chemotherapeutic drugs (etoposide, cisplatin, and vinblastin) (Jaboin et al. 2002; Scala et al. 1996). The survival of neuroblastoma tumor cells exposed to most chemotherapeutic drugs is mainly mediated by activation of the BDNF/TrkB

via activation of PI-3kinase/Akt and inactivation of GSK-38(Jaboin et al. 2002; Li et al. 2005, 2007a). But for some chemotherapeutic drug (paclitaxel), the survival of neuroblastoma cells is mediated by activation of MAPK pathway and inhibition of Bim (Li et al. 2007b). Such studies also demonstrated that the effects of chemotherapeutic agents are attenuated even if a tumor cell with high levels of TrkB expression is in an environment with low levels of BDNF. More importantly, if tumor cells express low levels of TrkB receptor, and are in a BDNF-rich environment, the effects of cytotoxic agents were decreased (Jaboin et al. 2002). These findings indicate that sites of relatively high levels of TrkB ligands (BDNF and NT-4) may serve as sanctuary sites for TrkB-expressing tumor cells. BDNF also stimulates tumor cell disaggregation and increases the ability of TrkB-expressing neuroblastoma tumor cells to degrade and invade through extracellular matrix proteins. The BDNF/TrkB-mediated invasiveness in neuroblastoma cells may be due to upregulation of Hepatocyte Growth Factor (HGF), originally described as Scatter factor and its receptor, c-met. The siRNA-mediated silencing of c-met or antibodies to HGF diminishes BDNF/TrkB-induced invasiveness. Whether this is the generalized mechanism by which TrkB mediates motility/invasiveness awaits evaluation in other model systems. Additionally, BDNF activation of TrkB increases tumor cell production of VEGF and this is enhanced in hypoxic conditions (Nakamura et al. 2006). Thus, the expression of both BDNF and full-length TrkB may represent an autocrine or paracrine survival pathway that is important for the aggressive behavior of some neuroblastoma (Acheson et al. 1995).

TrkC was first reported to be expressed predominantly in biologically favorable neuroblastomas (Ryden et al. 1996). Full-length TrkC is expressed in about 25 % of primary neuroblastomas, and these tumors essentially represent a subset of the TrkA-expressing tumors (Svensson et al. 1997; Yamashiro et al. 1996). So TrkC expression in neuroblastoma has been associated with good prognosis. The TrkC ligand NT-3 is recently reported to be overexpressed in 38 % of stage 4 neuroblastoma and also in a fraction of neuroblastoma cell lines mainly derived from stage 4 neuroblastoma tumor materials. Tumors with high NT-3 levels showed a high NT-3/TrkC ratio (Bouzas-Rodriguez et al. 2010). This observation is consistent with the theories of TrkC being a dependence receptor, which can initiate two completely opposite signaling pathways, depending on ligand availability (Tauszig-Delamasure et al. 2007). In the presence of ligand, a positive differentiation, guidance, or survival signal is transduced, while in the absence of ligand, receptors mediate an active process of apoptotic cell death. TrkA may also function as a dependence receptor (Nikoletopoulou et al. 2010). So autocrine NT-3 expression in stage 4 neuroblastoma may be a mechanism developed by a large fraction of tumor cells to bypass TrkC-induced cell death that would occur in regions of limited NT-3 concentrations.

**Medulloblastoma** is a central nervous system tumor which arises from the external granule cell layer of the cerebellum. TrkC expression was detected in medulloblastoma, and elevated TrkC expression is associated with favorable clinical outcome (Segal et al. 1994). Studies show that overexpression of TrkC inhibits the growth of intracerebral xenografts of a medulloblastoma cell line in nude mice

(Muragaki et al. 1997). NT-3 mRNA was also detected in medulloblastoma specimens, and in vitro studies find that adding NT-3 to TrkC-expressing medulloblastoma cells induces apoptosis as well as activation of multiple parallel signaling pathways, like PI3K, ERK, JNK/SAPK, p38MAPK. Of these, p38MAPK activity appears necessary for NT-3-/TrkC-induced apoptosis, based on experiments using pharmacological inhibitors. BDNF and TrkB were also detected in some medulloblastoma specimens. Treatment with BDNF of TrkB-expressing medulloblastoma cells significantly inhibited the cell viability in vitro. Besides, NGF treatment of TrkA-expressing medulloblastoma cells results in massive apoptosis (Muragaki et al. 1997), and this NGF/TrkA-induced apoptosis is mediated through Ras/Raf signaling pathway. Thus, activation of the neurotrophin/Trk signaling pathways plays a pro-apoptotic role in medulloblastoma.

## 5.2 Neurotrophins and Their Receptors in Nonneuronal Origin Cancers

#### 5.2.1 Colorectal Cancer

In colorectal cancer, BDNF and TrkB are overexpressed in the patients' tumors relative to levels in their adjacent normal tissues and this is more pronounced in advanced stages of colorectal cancer (Akil et al. 2011; Brunetto de Farias et al. 2010; Yu et al. 2010). In colorectal cancer cell lines, TrkB protein and BDNF mRNA expression are detected under basal (10 % serum) culture conditions, and BDNF mRNA levels and mature BDNF protein levels increase under stress condition (serum starvation) (Akil et al. 2011). Furthermore, in basal culture conditions, TrkB and BDNF have a cytosolic location in colorectal cell lines, but serum starvation induces a relocation of TrkB to the cell membrane and a colocalization of TrkB and BDNF on the membrane. BDNF induces cell proliferation and stimulates a TrkB-dependent anti-apoptotic effect under serum starvation condition. However, pro-BDNF, also detected in colorectal cancer cells, is co-expressed with p75NTR and sortillin at the cell membrane. In contrast to BDNF, exogenous pro-BDNF-induced colorectal cancer cell apoptosis, which suggests that a counterbalance mechanism is involved in the control of colorectal cancer cell survival (Akil et al. 2011). In human colorectal tumors, mutations in the TrkB kinase domain at two different sites (T695I, 751 N) have been found. The mutated TrkB displayed reduced activity compared to that of wild-type TrkB that indicated no gain-of-function of these mutations (Geiger et al. 2011). It is not known how these mutated TrkB receptors affect colorectal tumor cell biology. TrkA and TrkC were not detected in tumor tissues or in cell lines (Akil et al. 2011).

#### 5.2.2 Breast Cancer

High levels of TrkA and phospho-TrkA are expressed in breast cancer tumors (compared to normal breast tissues) and in breast cancer cells (Davidson et al. 2004; Lagadec et al. 2009). Overexpression of TrkA induces constitutive activation of its tyrosine kinase activity. This promotes cell growth, migration, and

invasion in vitro via the activation of PI3K/Akt and MAPK signaling pathways which are required for the maintenance of a more aggressive cellular phenotype. TrkA overexpression enhances tumor growth, angiogenesis, and metastasis of breast cancer xenografts in immunodeficient mice (Lagadec et al. 2009).

NGF expression is detected in most breast tumors as well as in breast cancer cell lines, but no direct relationship between NGF overexpression and any known prognostic factor has been observed. Inhibition of NGF with neutralizing antibodies, or small interfering RNA, strongly reduces angiogenesis and tumor xenograft development in immunodeficient mice (Adriaenssens et al. 2008). Since NGF activation of TrkA has been shown to increase VEGF production by neuroblastoma tumor cells (Nakamura et al. 2011), it is possible that the decreased angiogenesis results from decreased tumor cell production of VEGF when NGF is inhibited. The mitogenic effect of NGF on breast cancer cells requires the tyrosine kinase activity of TrkA as well as the mitogen-activated protein kinase (MAPK) cascade, but is independent of p75NTR. In contrast, the anti-apoptotic effect of NGF is not dependent on TrkA but rather p75NTR and the activation of the transcription factor NF- $\kappa$ B (Descamps et al. 2001).

Truncated TrkB (TrkB.T1) and p75NTR, as well as BDNF and NT-4/5, are detected in breast cancer specimens and breast cancer cells. Full-length TrkB (TrkB.FL) has only rarely been reported (Vanhecke et al. 2011). Blocking endogenous NT-4/5 and BDNF using neutralizing anti-NT-4/5 or anti-BDNF antibodies inhibits the cell survival. Adding exogenous BDNF or NT-4/5 has an anti-apoptotic effect in breast cancer cells, but little effect on cell proliferation and invasion. Targeted decrease in p75NTR or TrkB-T1 using siRNAs abolishes the NT-4/5 and BDNF anti-apoptotic effect. Moreover, the Trk pharmacological inhibitor K252a has no impact on the BDNF- and NT-4/5-mediated survival, suggesting the involvement of p75NTR and TrkB-T1, and not Trk kinase activity. Treatment of mice bearing breast cancer xenografts with anti-BDNF, anti-NT-4/5, anti-p75NTR, or anti-TrkB-T1 inhibits tumor growth and this is accompanied by an increase in cell apoptosis and no changes in cell proliferation. A study showed that higher expression of BDNF is significantly associated with unfavorable pathologic parameters and poor prognosis (Patani et al. 2011). This indicates that an autocrine loop of BDNF and NT-4/5 stimulates breast cancer cell survival.

TrkC is also detected in breast cancers, and there is an inverse relationship between tumor grade and TrkC expression. The expression of TrkC is higher in Grade I tumors that have a good prognosis compared to that in Grade III tumors which have a poor prognosis. Rise in TrkC expression could indicate good prognosis (Blasco-Gutierrez et al. 2007).

#### 5.2.3 Prostate Cancer

TrkA receptor is expressed in normal prostate, and expression is retained or significantly overexpressed in prostate cancer (MacGrogan et al. 1992; Mahadeo et al. 1994). TrkB and TrkC receptors are only expressed in malignant prostate cells. p75NTR is expressed in normal human prostate epithelial cells, and its expression is progressively lost during progression of prostate cancer. There is an

inverse relationship between low p75NTR expression and high prostate cancer growth and metastasis. So p75NTR may function as a prostate tumor-suppressor and metastasis-suppressor gene (Krygier and Djakiew 2002). TrkA activation increases cell proliferation, while p75NTR expression is associated with apoptosis. So an imbalance between p75NTR and TrkA-mediated signals may be involved in the progression of prostate cancer through increased proliferation and reduced apoptosis.

Human prostate is the second most abundant source of NGF after central nervous system. NGF is detected in prostate cancer specimens and cell lines. NGF secreted by the prostate stromal cells stimulates the growth of prostate cancer cells, and some specific prostate tumor cell lines secrete NGF in an autocrine manner (Djakiew 2000). Thus, NGF contributes to prostate cancer growth in both a paracrine and an autocrine manner.

#### 5.2.4 Lung Cancer

Trk receptors but not p75NTR are readily detected in multiple types of human lung cancer (Koizumi et al. 1998; Ricci et al. 2001). High levels of TrkA are observed in adenocarcinomas and squamous cell carcinomas, while TrkB is detected primarily in small cell lung cancers (SCLC). Some tumors also express neurotrophin ligands. In particular, NGF is expressed in a high proportion of bronchioloalveolar and squamous cell carcinomas, while BDNF is expressed in SCLC. Approximately one third of bronchioloalveolar carcinomas express both NGF and TrkA, an observation that raises the possibility of autocrine effects of neurotrophins in this cancer type. Such autocrine loops may also exist as BDNF and TrkB expression is co-expressed in adenocarcinomas, squamous cell carcinomas, and small cell lung cancers. NT-3 and its corresponding TrkC receptor are found in a small number of squamous cell carcinomas (Ricci et al. 2001).

Studies with lung tumor cell lines indicate that NGF inhibits the growth of SCLC cells, and this is accompanied by an upregulation of TrkA, p75NTR, and NGF expression. NGF markedly inhibits cellular proliferation, colony formation, and invasiveness in vitro. NGF-treated cell lines do not form tumors in nude mice, whereas the untreated cell lines are highly tumorigenic. Administration of NGF to mice bearing established xenografts inhibits tumor growth.

In non-small cell lung cancers (NSCLC), TrkB enhances cell migration, invasion, and transformation (Harada et al. 2011), and a higher expression of TrkB in NSCLC primary tumors is closely correlated with lymph node metastasis (Zhang et al. 2010).

#### 6 Clinical-Translational Advances

After the discovery of the involvement of neurotrophins and their receptors in the biological and clinical behaviors of cancers, it was appreciated that inhibition of neurotrophins, their receptors, or their targets might be beneficial in the clinical oncology setting. One well-studied strategy targets the Trk tyrosine kinase domain. Since there is a high degree of homology among Trk tyrosine kinase domains, there

are currently no inhibitors reported that selectively block TrkA or TrkB kinase activity. The earliest Trk inhibitors to reach clinical trials were Cephalon's CEP-751 and CEP-701 (lestaurtinib). These compounds were modeled on the indolocarbazole K252a from Nocardiopsis species and were relatively nonselective in inhibiting Trks, as well as Flt3, JAK2, and protein kinase C (PKC). CEP-751 and CEP-701 reduced tumor burden in xenograft models of prostate and pancreatic cancer, neuroblastoma, medulloblastoma, and medullary thyroid carcinoma (Camoratto et al. 1997; Evans et al. 1999, 2001; George et al. 1999; Miknyoczki et al. 1999). In prostate cancers, CEP-701 also inhibits the metastasis of both primary and metastatic sites of prostate cancer. A phase 1 study of CEP-701 established the maximum tolerated dose and identified that toxicities such as nausea, gastrointestinal events, and dyspepsia were relatively well tolerated (Marshall et al. 2005). Recently, a phase 1 study indicated that CEP-701 was well tolerated in patients with refractory neuroblastoma, and a dose level sufficient to inhibit TrkB activity was established (Minturn et al. 2011). CEP-701 phase 2 studies are in progress. Given the important role Trk signaling plays in normal neurons, it is encouraging that neural toxicity has not been a dose-limiting toxicity.

Another series of Trk inhibitors came from the optimization of 4aminopyrazolylpyrimidines (AstraZeneca, Inc). One such compound, AZ-23, inhibits TrkA/B in the nanomolar range and has reasonable pharmacodynamics and oral bioavailability in toxicology studies. Phase 1 testing in adults was initiated but recently halted as a result of poor pharmacodynamic properties (Wang et al. 2008). Study on another compound named AZ623 showed that AZ623 inhibits BDNF-mediated signaling and neuroblastoma cell proliferation in vitro, as well as also inhibits the growth of human neuroblastoma xenograft tumors in vivo (Zage et al. 2011). Other congeners of 4-aminopyrazolylpyrimidines remain poised for clinical development.

Another strategy suggested by preclinical modeling is to use Trk kinase inhibitors in combination with standard chemotherapy. In a preclinical neuroblastoma xenograft model, CEP-701 (or the active precursor CEP-751) significantly enhanced the activity of cyclophosphamide, as well as other cytotoxic drug combinations currently used in the treatment of high-risk neuroblastoma (Evans et al. 1999). A clinical trial of CEP-701 with gemcitabine was undertaken assessing drug interactions in patients with advanced pancreatic cancer (Chan et al. 2008). The combination was well tolerated with no unexpected toxicities, but because the trial was limited and not designed to assess efficacy, further studies are planned. A study in a combination of AZ623 with topotecan, a topoisomerase inhibitor that is currently used for neuroblastoma treatment, indicated that a combination of AZ623 and topotecan has much higher effect on the inhibition of xenograft tumor growth compared to either individual agent, and inhibits the regrowth of tumors after cessation of the treatment where either individual agent alone is ineffective (Zage et al. 2011).

Additional therapeutic options to block the effects of activated Trk receptors can be developed by moving downstream and targeting the key intracellular mediators of activated Trk's pro-survival and prometastatic signals, PI3K and Akt. Pharmacologic inhibitors of these "nodal" kinases have been extensively pursued. Perifosine is an oral alkyl-phospholipid Akt inhibitor. Unlike most kinase inhibitors, which target the adenosine triphosphate-binding region, perifosine targets the pleckstrin homology domain of Akt. Phase I and II clinical trials have indicated that the side effect of perifosine is limited to dose-limiting gastrointestinal toxicity, such as nausea, diarrhea, fatigue, and dehydration, which can be ameliorated with the use of prophylactic medicine and well tolerated (Gills and Dennis 2009). As Akt is a key downstream target in PI3K pathway initiated by many growth factors and their receptors, including neurotrophins/Trk, perifosine as an Akt inhibitor has been evaluated in many types of tumors with elevated activity of Akt, and has showed antitumor growth effect either alone or in combination with chemotherapeutic drugs/other small molecular inhibitors. The role of perifosine in tumors with BDNF/TrkB-initiating activation of Akt has been evaluated in neuroblastoma. Perifosine alone inhibits tumor growth, but is most effective in combination with cytotoxic drugs such as etoposide. Combinations of Akt inhibition and etoposide cause tumor regression of TrkB-expressing neuroblastoma tumor xenografts (Li et al. 2011). Strategies targeting Akt may offer a broader spectrum of inhibitory activity than Trk kinase-targeted agents and may avoid resistance due to feedback suppression that has been seen when targeting some receptors.

# 7 Conclusion

Neurotrophins and their receptors were originally found to play important roles in the survival, proliferation, differentiation, apoptosis, and synaptic plasticity of neurons, with aberrant expression being associated with neurodegenerative diseases. However, increasingly studies have indicated that the neurotrophin/neurotrophin receptors play an important role during the genesis of tumors where oncogenic events may subvert the normal functions of neurotrophins and their receptors and influence biologic features of tumors that may impact tumor cell aggressiveness. Neurotrophins activate different signaling pathways based on binding to their Trk receptors or p75NTR receptor, and they may exert pro- or anti-tumorigenic activity, depending on the activation status of other cellular receptors and the presence or absence of ligands in the environment. For most types of cancers, expression of TrkA or TrkC in tumor tissues is associated with good prognosis, such as neuroblastoma, medulloblastoma, lung cancer; expression of TrkB and/or its ligand BDNF is associated with poor prognosis, such as neuroblastoma and colorectal cancer. One exception is breast cancer where overexpression of TrkA and NGF in breast cancer cells stimulates cell proliferation and invasion but not cell differentiation or apoptosis. These findings provide support for the development of small molecular inhibitors that target on Trk or the downstream targets in the neurotrophin/neurotrophin receptor-initiating signaling pathways in the treatment of cancers. The preclinical and clinical evidence of efficacy as a single agent or in combination with chemotherapy, as well as the welltolerated limited toxicities, suggests the usage of these small molecular inhibitors as a potential strategy for the treatment of cancers.

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# **NGF and Immune Regulation**

# Luisa Bracci-Laudiero and Luigi Manni

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#### Abstract

For some time after its discovery, the action of nerve growth factor (NGF) was considered restricted to the nervous system. Instead, a variety of experimental data indicate that NGF can influence the activity of both the nervous and immune systems. This should not be surprising since these two systems are responsible both for maintaining homeostasis and for adapting the body to the environment. To orchestrate strictly integrated responses, they need to have close anatomical connections and to share common chemical signals and specific receptors. The well-known effects of NGF on peripheral neuron survival and maintenance and dynamic control by NGF of innervation and neuropeptide synthesis, together with its direct effects on immune cell functions, indicate that NGF has a key role in the complex network of bidirectional signals between the nervous and immune systems. NGF receptors are expressed in immune organs and cell populations, allowing NGF to modulate cell differentiation and regulate immune response. NGF concentrations in tissues change during inflammation, and inflammatory mediators induce NGF

L. Bracci-Laudiero (🖂) • L. Manni (🖂)

Institute of Translational Pharmacology, National Research Council of Italy (CNR), Rome, Italy e-mail: luisa.braccilaudiero@opbg.net; luisa.braccilaudiero@ift.cnr.it; luigi.manni@ift.cnr.it

synthesis in a variety of cell types. As a growing number of studies have shown, an enhanced production of NGF characterizes inflamed tissues of patients with inflammatory and autoimmune diseases. Unfortunately, although the dynamic regulation of NGF synthesis seems to be a common feature of chronic inflammatory diseases, the reasons why NGF concentrations are enhanced and how this can affect inflammatory responses and the course of the diseases are far from being understood.

List of Abbreviations				
BBB	Blood-brain barrier			
CGRP	Calcitonin gene-related peptide			
CHS	Contact hypersensitivity			
CIPA	Congenital insensitivity to pain with anhidrosis			
CNS	Central nervous system			
COX2	Cyclooxygenase 2			
CSF	Cerebrospinal fluid			
DRG	Dorsal root ganglion			
EAE	Experimental autoimmune encephalomyelitis			
Gal	Galanin			
GM-CSF	Granulocyte-macrophage colony-stimulating factor			
IFN	Interferon			
IL	Interleukin			
LPS	Lipopolysaccharide			
MAPK	Mitogen-activated protein kinase			
MPB	Myelin basic protein			
NGF	Nerve growth factor			
NO	Nitric oxide			
NPY	Neuropeptide Y			
NT	Neurotrophin			
NTR	Neurotrophin receptor			
PI3K	Phosphatidylinositol 3-kinase			
PNS	Peripheral nervous system			
SP	Substance P			
TH	Tyrosine hydroxylase			
TNF	Tumor necrosis factor			
TrkA	Tropomyosin kinase receptor A			
UV	Ultraviolet			
VIP	Vasoactive intestinal peptide			

# 1 Introduction

The neurotrophin (NT) family of proteins is responsible for the growth and survival of neurons during development and for the maintenance and function of adult neurons (Rush et al. 1997; Ernfors 2001; McAllister 2001; Poo 2001). NTs also promote

regeneration of damaged axons after various peripheral and central nervous system injuries (Olson 1993; Connor and Dragunow 1998; Terenghi 1999; Chao et al. 2006).

Nerve growth factor (NGF) was the first NT to be discovered (Levi-Montalcini 1987). It is essential for the development and maintenance of neurons in the peripheral nervous system (PNS) and for the functional integrity of cholinergic neurons in the central nervous system (CNS) (Whittemore et al. 1987). For over 35 years, NGF has been considered a very powerful and selective growth factor for sympathetic and sensory neurons and for cells derived from the neuronal crest (Alleva et al. 1993; Rush et al. 1997; Cowan 2001). In the CNS, the phenotypic maintenance of cholinergic neurons is dependent on NGF, which is produced mainly in the cortex, the hippocampus, and the pituitary gland, although significant quantities of this NT are also produced in other brain areas, including the basal ganglia, thalamus, hypothalamus, and spinal cord, and in the retina (McAllister 2001).

NGF exerts its biological action by challenging the specific receptor tropomyosin kinase receptor A (TrkA), a typical tyrosine kinase receptor (Huang and Reichardt 2003). The major cytosolic/endosomal pathways activated by TrkA are Ras-mitogen-activated protein kinase (MAPK), extracellular signal-regulated kinase (ERK), phosphatidylinositol 3-kinase (PI3K)-Akt, and PLC- $\gamma$  (Klesse and Parada 1999; Chao et al. 2006; Reichardt 2006). NGF also binds to and activates the nonselective p75 pan-neurotrophin receptor (p75<sup>NTR</sup>). This receptor is a transmembrane glycoprotein that regulates signaling through TrkA (Friedman and Greene 1999; Schor 2005; Reichardt 2006); the binding of NGF to p75<sup>NTR</sup> activates additional signaling pathways that, in the absence of co-expressed TrkA, may signal cell death via apoptosis (Friedman and Greene 1999; Miller and Kaplan 2001; Schor 2005). Signaling pathways activated by p75<sup>NTR</sup> are the Jun kinase signaling cascade, NF-κB, and ceramide generation (Reichardt 2006).

The pioneering studies of Levi-Montalcini showed that, during development, limited amounts of NGF produced in target organs are responsible for the proliferation, differentiation, and survival of peripheral autonomic and sensory neurons (Levi-Montalcini 1987). Moreover, neutralizing endogenous NGF with anti-NGF antibodies during fetal life causes permanent damage to the sympathetic ganglia and a marked decrease in sensory neuron numbers (Levi-Montalcini 1987). In adult life, while sympathetic neurons continue to depend on NGF for their survival (Ruit et al. 1990), there is a change in the role of NGF in the sensory neurons. NGF is no longer essential for their survival but remains crucial for the maintenance of a differentiated phenotype (McMahon et al. 1995; Verge et al. 1995). In these mature sensory neurons, as well as in sympathetic neurons, NGF dynamically controls neurotransmitter and neuropeptide synthesis. In sympathetic neurons, the levels of norepinephrine are regulated by NGF through the selective induction of tyrosine hydroxylase (TH) (Otten et al. 1977). In dorsal root ganglia (DRG), the expression of neuropeptides such as substance P (SP) and calcitonin gene-related peptide (CGRP) in primary sensory neurons is under NGF control (Lindsay and Harmar 1989), and in vivo deprivation of NGF, following nerve transection or anti-NGF treatment, causes a marked decrease in SP and CGRP synthesis (Verge et al. 1995).

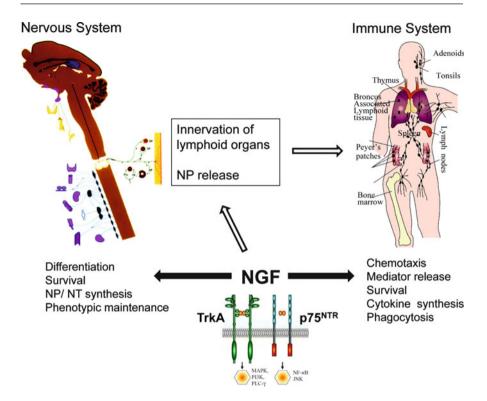
The NGF supplied from the innervation field influences the neuronal plasticity that allows the adult nervous system to modify its structure and functions in response to stimuli. Indeed, the constitutive synthesis of NGF in adult tissues correlates with the phenotypic features of PNS neurons, such as innervation density, cell body size, axonal terminal sprouting, dendritic arborization, and induction or inhibition of neuropeptides and neurotransmitters or transmitter-producing enzymes (Otten et al. 1977; Lindsay and Harmar 1989; Mearow and Kril 1995; Verge et al. 1995).

The spectrum of NGF action was once considered limited to functions related to specific CNS and PNS neuronal classes; however, more recent studies have revealed that the effects of NGF also extend to several kinds of nonneuronal cells. Indeed, NGF can influence the functioning of the immune-hematopoietic system (Aloe et al. 1997a; Bonini et al. 1999; Aloe 2001) and the neuro-endocrine system (Aloe et al. 1997a). Thus, the current view is that *NGF can affect the maintenance of a balanced interplay between the nervous, endocrine, and immune systems* (Alleva et al. 1993; Aloe et al. 1999, 2002) (Fig. 1).

#### 2 NGF and the Immune System

## 2.1 NGF Influences Development and Differentiation of the Immune System

The release of neuropeptides from sensory nerve endings or of neurotransmitters from autonomic terminals represents a direct means for the nervous system to regulate immune response. Indeed, neurotransmitters and/or neuropeptide receptors are expressed by immune organs and cells, and their challenge has immunemodulatory value (Elenkov 2008; Tausk et al. 2008). NGF regulates neuropeptides and neurotransmitters in peripheral neurons by acting on their promoters or on neurotransmitter-producing enzymes (Otten et al. 1977; Lindsay and Harmar 1989; Verge et al. 1995). Thus, a possible mechanism by which NGF can affect the immune response is through the regulation of neurotransmitter and neuropeptide production. But NGF can also influence immune cell functions directly, as both NGF and NGF receptors are expressed in lymphoid organs and in purified immune cell populations. The expression of NGF receptors has been found in primary (thymus, bursa of Fabricius, bone marrow) and secondary (spleen, tonsils, lymph nodes) lymphoid organs (Laurenzi et al. 1994; Ciriaco et al. 1996; Garcia-Suarez et al. 2001), being raised during lymphocyte precursor migration but declining during postnatal life. NGF is produced in primary lymphoid organs (Laurenzi et al. 1994; Aloe et al. 1997b), and it is possible that localized synthesis of NGF may contribute to create the microenvironment for the differentiation of hematopoietic stem cells (Pezzati et al. 1992; Cattoretti et al. 1993). In vitro NGF stimulates the functional activity of thymic epithelial cells, inducing the expression of adhesion molecules important for thymocyte-thymic epithelia interaction, and upregulates the expression of thymopoietic factors such as IL-7, granulocytemacrophage colony-stimulating factor (GM-CSF), and stromal cell-derived



**Fig. 1** *Nervous-immune systems interplay.* The NGF is at the interface of the nervous and immune system interactions. By activation of the specific TrkA receptor and/or the pan-neurotrophin p75 receptor, it acts on nervous system (*left side*) regulating survival, differentiation, and phenotypic features such as the synthesis of neurotransmitters and neuropeptides or the shape of cells and terminals. NGF influences immune system development and activity directly, by challenging NGF receptors expressed on the surface of immune cells. NGF also indirectly modulates immune system functions by regulation of the activity of nerve terminals innervating lymphoid organs and of the synthesis and release of neurotransmitters and neuropeptides that in turn activate specific receptors expressed by immune cells/organs. Abbreviations: *NP* neuropeptides, *NT* neurotransmitters, *TrkA* tyrosine kinase A receptor, *p75<sup>NTR</sup>* p75 neurotrophin receptor

factor-1 (Lee et al. 2008). TrkA expression has been described in hematopoietic precursors (Chevalier et al. 1994) and NGF functions as a survival factor (Bracci-Laudiero et al. 1993b; Auffray et al. 1996) and a colony-stimulating factor (Matsuda et al. 1988, 1991; Welker et al. 2000) for myeloid progenitors. Human CD34+ hematopoietic stem cells are characterized by the concomitant expression of TrkA and production of their own NGF, which is maintained during lineage differentiation but declines with the acquisition of a mature phenotype (Bracci-Laudiero et al. 2003). Moreover, the changes in TrkA expression observed in immature cord blood cells and mature immune cells may indicate a different requirement for NGF in these cells that depends on the state of maturity and functional activity of the immune cell (Bracci-Laudiero et al. 2003) (Fig. 2).

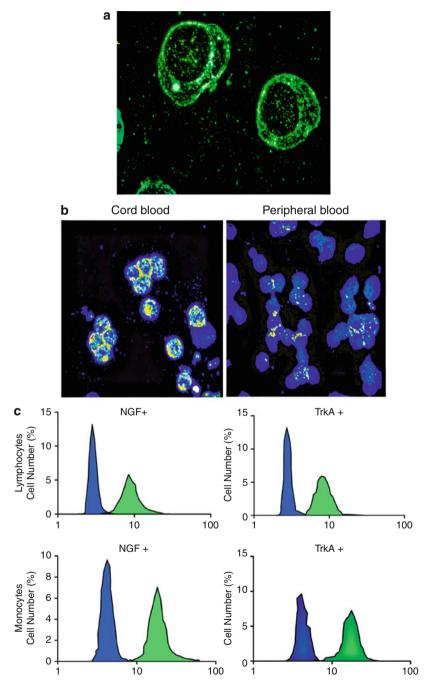
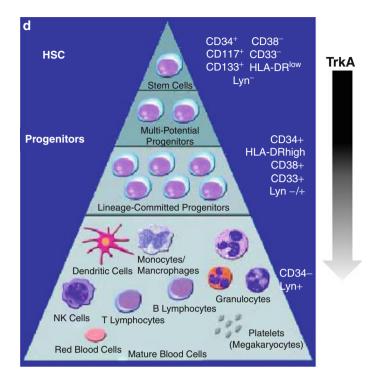


Fig. 2 (continued)



**Fig. 2** *NGF influences the differentiation of hemopoietic stem cell.* TrkA (**a**) and NGF (**b**) expression is detected by immunofluorescence in CD34+ stem cells purified from human umbilical cord blood (HCB). Flow cytometry analysis (**c**) reveals that NGF and TrkA are co-expressed in the lymphocytic and monocytic components of HCB. As a clue of the importance of NGF in the development of immune cell lineage, the expression of TrkA is decreasing on immune cell surface as they become more differentiated (**d**)

#### 2.2 NGF and Immune Cells

NGF receptor expression has been studied in purified immune cell populations. TrkA expression has been demonstrated in mononuclear cells (Thorpe et al. 1987a), thymocytes (Laurenzi et al. 1994), human B and T lymphocytes (Brodie and Gelfand 1992; Ehrhard et al. 1993; Torcia et al. 1996; Lambiase et al. 1997), monocytes (Ehrhard et al. 1993; Caroleo et al. 2001), mast cells (Nilsson et al. 1997), basophils (Burgi et al. 1996), and eosinophils (Noga et al. 2002). These cells have a basal expression of TrkA in the resting condition and upregulate it after antigenic or mitogenic stimulation, when functional activation is necessary (Ehrhard et al. 1993a, b; Torcia et al. 1996; Lambiase et al. 1997; Caroleo et al. 2001). The evidence that almost all immune cells express NGF receptors provided great impulse to studies on the influence of NGF on immune functions. The study demonstrating that the injection of NGF in newborn rats caused an increase in the number and size of mast cells (Aloe and Levi-Montalcini 1977) was the earliest

evidence of an effect of NGF on immune cells and outside the nervous system. After this, numerous studies have shown that NGF is able to exert a wide spectrum of effects on immune cells (Table 1). One of the effects of NGF that emerges as common to the greater part of cell populations analyzed is its dose-dependent influence on the survival of hematopoietic stem cells and mature, differentiated immune cells (Auffray et al. 1996; Bullock and Johnson 1996; Torcia et al. 1996; Bracci-Laudiero et al. 2003). The viability-sustaining activity of NGF has been attributed to its effect on the induction of anti-apoptotic proteins such as Bcl-2 (Bullock and Johnson 1996).

#### 2.2.1 Granulocytes and Mast Cells

In general, in vitro studies of granulocytes and mast cells have shown that NGF can potentiate their responses to inflammatory stimuli, but its effects differ depending on the cell type analyzed. In activated basophils, NGF enhances the synthesis of lipid mediators (leukotrienes) in response to inflammatory stimuli (Bischoff and Dahinden 1992; Takafuji et al. 1992) and, together with IL-3 and IL-33, acts as cofactor to induce the release of histamine and enhance the response to IgE in both basophils and mast cells (Mazurek et al. 1986; Horigome et al. 1993; Sin et al. 2001; Gibbs et al. 2005). In contrast to its effect on basophils, NGF tends to suppress leukotriene formation in eosinophils (Takafuji et al. 1992). However, NGF seems to stimulate the production of IL-4 (Noga et al. 2002); it influences the release of peroxidase in eosinophils and enhances cytotoxic activity (Hamada et al. 1996). It also enhances, in a dose-dependent way, the survival of eosinophils (Hamada et al. 1996) and neutrophils (Kannan et al. 1991) by suppressing apoptosis, an effect abolished by anti-NGF antibody treatment. In neutrophils, in addition to its survival effect, NGF enhances superoxide production after cell activation with zymosan and stimulates phagocytosis (Kannan et al. 1991).

As for granulocytes, a pro-survival effect of NGF has also been demonstrated in mast cells (Kawamoto et al. 1995; Bullock and Johnson 1996). NGF also influences mast cell activation in various stages of cell differentiation: it causes a dose-dependent upregulation of tryptase and IgE receptors in immature mast cells (Welker et al. 1998), while the addition of NGF after stimulation of mature mast cells with lysophosphatidylserine induces and enhances cyclooxygenase2 (COX2) expression and the generation of prostaglandin D2 (Murakami et al. 1997).

Several studies have shown that NGF is a chemoattractant factor for both granulocytes (Gee et al. 1983; Boyle et al. 1985) and mast cells (Sawada et al. 2000). The migration of eosinophils towards the site of inflammation in the airways is influenced by the presence of NGF, and this effect seems to be regulated by neuropeptides (Quarcoo et al. 2004). Circulating neutrophils also seem to be involved in NGF-induced hyperalgesia in the skin of rat paw (Bennett et al. 1998). Indeed, local administration of NGF in rat skin induced hyperalgesia and neutrophil accumulation, an effect that appears to be mediated by the induction by NGF of adhesion molecules (i.e., ICAM-1) in endothelial cells (Foster et al. 2003). The use of anti-ICAM-1 antibodies blocked neutrophil accumulation in

	Cell type	NGF-/TrkA-regulated functions	References
۲	Hematopoietic stem cell	Survival, differentiation	Matsuda et al. (1991), Bischoff and Dahinden (1992), Takafuji et al. (1992), Cattoretti et al. (1993), Chevalier et al. (1994), Auffray et al. (1996), Bracci-Laudiero et al. (2003), Sato et al. (2004)
Ø	Eosinophils	Differentiation, mediator release	Takafuji et al. (1992), Hamada et al. (1996), Noga et al. (2002), Quarcoo et al. (2004)
6	Basophils	Differentiation, mediator release	Bischoff and Dahinden (1992), Takafuji et al. (1992), Burgi et al. (1996), Sin et al. (2001), Gibbs et al. (2005)
	Neutrophils	Chemotaxis, phagocytosis, mediator release, survival	Gee et al. (1983), Boyle et al. (1985), Kannan et al. (1991), Bennett et al. (1998), Beigelman et al. (2009)
Ø	Mast cells	Degranulation, mediator release, survival	Mazurek et al. (1986), Matsuda et al. (1991), Kawamoto et al. (1995), Lambiase et al. (1995), Bullock and Johnson (1996), Lowe et al. (1997), Murakami et al. (1997), Nilsson et al. (1997), Welker et al. (1998), di Mola et al. (2000), Sawada et al. (2000), Welker et al. (2000), Xiang and Nilsson (2000), Stanzel et al. (2008)
	B lymphocytes	Proliferation, differentiation, IgG synthesis, survival, CGRP synthesis	Otten et al. (1989), Kimata et al. (1991), Brodie and Gelfand (1992), Franklin et al. (1995), Brodie et al. (1996), Melamed et al. (1996), Torcia et al. (1996), Bracci-Laudiero et al. (2002), Coppola et al. (2004), Sato et al. (2004), Abram et al. (2009)
	T lymphocytes	Proliferation, NPY synthesis	Thorpe et al. (1987b), Ehrhard et al. (1993), Bracci-Laudiero et al. (1996a), Lambiase et al. (1997)
	Monocytes	Oxidative burst, cytokine release, survival, CGRP release	Ehrhard et al. (1993), Susaki et al. (1996), la Sala et al. (2000), Barouch et al. (2001), Caroleo et al. (2001), Bracci-Laudiero et al. (2002), Bracci-Laudiero et al. (2005), Noga et al. (2007), Samah et al. (2009), Ma et al. (2010)

## Table 1 Summary of main effects of NGF on immune cells

NGF-treated skin, suggesting that NGF has primarily an activating effect on endothelial cells, resulting in the accumulation of neutrophils (Foster et al. 2003). Abnormally decreased migration of neutrophils has been reported in patients with congenital insensitivity to pain with anhidrosis (CIPA), a rare genetic disease characterized by mutation of the TrkA gene (Beigelman et al. 2009). All these data suggest that the presence of *NGF is pivotal to cell recruitment at the inflammation site*, though the mechanisms and mediators involved still remain to be clarified.

## 2.2.2 B and T Lymphocytes

Experimental evidence produced in the 1980s indicated that NGF receptors are expressed on lymphocytes and that NGF is able to potentiate the proliferative response of B- and T-cells to mitogens (Thorpe and Perez-Polo 1987; Thorpe et al. 1987a, b). The addition of NGF to culture media induces interleukin-2 receptors in mature lymphoid cells, and the proliferative response is augmented in B-cells when NGF is added together with IL-2 (Brodie and Gelfand 1992). Although CD4+ lymphocytes also express TrkA receptors (Ehrhard et al. 1993a), few data on these cells are available, while more numerous studies have focused on B lymphocytes. B-cells express TrkA (Torcia et al. 1996), and the binding of NGF to this receptor activates intracellular pathways and nuclear factors in a manner similar to that described in neurons (Franklin et al. 1995; Melamed et al. 1996). Abolition of TrkA function in knockout mice models (Coppola et al. 2004) or in CIPA patients (Sato et al. 2004) results in altered differentiation and survival of B lymphocytes. Moreover, NGF can influence the production of IgM and IgG (especially IgG4) (Otten et al. 1989; Kimata et al. 1991). Overall these results support the conclusion that NGF stimulates B-cell immune responses.

However, other reports describe evident contradictions to those cited above, with findings showing that NGF is a survival factor for memory B-cells (Torcia et al. 1996), promotes the differentiation of B-cells into immunoglobulin-secreting plasma cells (Otten et al. 1989), and influences plasma cell survival (Abram et al. 2009). Activation of B-cells with anti-CD40 plus NGF decreases instead of increasing immunoglobulin secretion in a dose-dependent manner (Brodie and Gelfand 1994; Brodie et al. 1996). Moreover, the finding that TrkA knockout mice have increased IgM, IgG1, and IgG2 levels (Coppola et al. 2004) suggests that a clear understanding of the physiological effects of NGF on B-cells is still lacking.

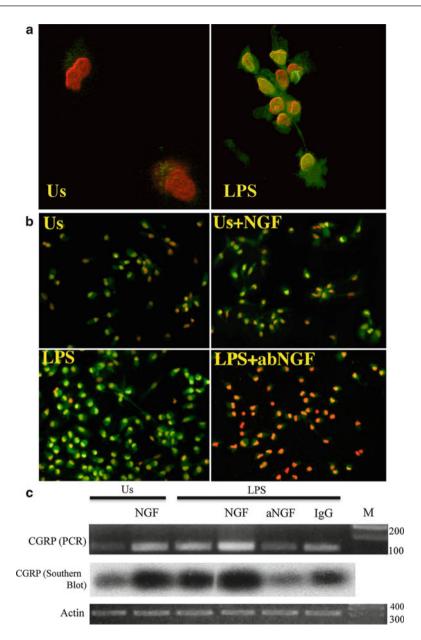
## 2.2.3 Monocytes/Macrophages and Dendritic Cells

Monocytes and macrophages express TrkA and  $p75^{NTR}$ , while differentiated dendritic cells lose TrkA and maintain  $p75^{NTR}$  expression only (Ehrhard et al. 1993b; la Sala et al. 2000; Caroleo et al. 2001). Inflammatory stimuli enhance (Ehrhard et al. 1993b; Caroleo et al. 2001) or induce a reexpression (Noga et al. 2007) of TrkA in all these cells. NGF protects monocytes from apoptosis

induced by UVB radiation or gliotoxin, regulating the expression of anti-apoptotic proteins Bcl-2, Bcl-xl, and Bfl-1 (la Sala et al. 2000). In vitro migratory properties are also affected by NGF, which increases CXCR4 expression and chemotactic response to suboptimal CXCL-12 concentrations in macrophages (Samah et al. 2009). Inflammatory macrophages isolated from the peritoneum after stimulation and then treated with NGF show increased phagocytosis, enhanced parasite-killing activity, and increased production of IL-1beta (Susaki et al. 1996). NGF can also induce TNF-alpha secretion and increase nitric oxide (NO) production in macrophages, this latter effect being strongly amplified by pretreatment with interferon-gamma (Barouch et al. 2001). NGF promotes LPS-induced maturation and secretion of inflammatory cytokines in dendritic cells (Noga et al. 2007). Dendritic cells derived from allergic donors and stimulated with LPS and CD40 ligand increase IL-6 production when challenged with NGF, while the effect of NGF on dendritic cells generated from healthy donors is to induce more IL-10 production (Noga et al. 2007). Consistent with this finding is the fact that NGF deprivation in LPS-treated monocytes results in a significant decrease in IL-10 synthesis (Bracci-Laudiero et al. 2005).

## 2.3 NGF Induces Neuropeptide Production in Immune Cells

Besides its ability to induce inflammatory mediators and cytokines, NGF can also modulate immune response by regulating the synthesis of neuropeptides in immune cells. Similarly to its effect on neuronal cells, NGF induces NPY expression in T lymphocytes (Bracci-Laudiero et al. 1996a) and regulates the expression and release of CGRP in human B lymphocytes (Bracci-Laudiero et al. 2002) and monocytes (Bracci-Laudiero et al. 2005; Ma et al. 2010) and macrophages. By inducing CGRP and IL-10 production, NGF can reduce the antigen-presenting capacity and co-stimulatory function of monocytes and may contribute to the downregulation of T-cell responses. Indeed, using CGRP receptor antagonists LPS-treated monocytes, the expression of membrane in molecules involved in antigen presentation, such as HLA-DR and CD86, is increased (Bracci-Laudiero et al. 2005). These data suggest that at least some of the NGF effects described to date are not directly dependent on NGF itself but rather are regulated by intermediate mediators whose synthesis is under NGF control. Neuropeptides, such as SP, CGRP, NPY, and VIP, have all been indicated as possible mediators of NGF effects on immune system response, in regulating inflammatory and anti-inflammatory responses and in maintaining homeostasis (Brain 1997; Delgado et al. 2004; O'Connor et al. 2004; Prod'homme et al. 2006). The point that deserves underlining is that NGF can act on immune cells, stimulating neuropeptide synthesis in both the neuronal and *immune systems* (Fig. 3).



**Fig. 3** *NGF induces CGRP in cultured macrophages.* Purified macrophages express calcitonin gene-related peptide (CGRP), as revealed by green fluorescence, if challenged with lipopolysac-charide (LPS) (a). Neutralizing NGF by specific anti-NGF antibody counteracted the CGRP expression in macrophages activated by exposure to LPS (b). Gene expression analysis (c) by PCR and Southern blot reveals that both unstimulated (Us) and LPS-challenged macrophages increase CGRP gene expression when exposed to NGF in culture, while anti-NGF decreases CGRP mRNA in stimulated cells

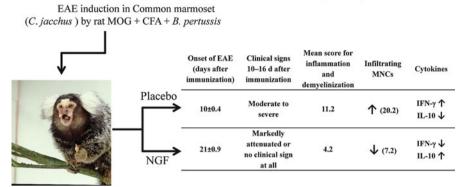
## **3** NGF and Inflammation

#### 3.1 In Vivo Evidence and Experimental Models

An increase in NGF in tissues at the site of inflammation is a characteristic of several inflammatory diseases. Increased NGF concentrations, which closely follow the course of the disease, were initially found in the cerebrospinal fluid of multiple sclerosis patients (Bracci-Laudiero et al. 1992), in the synovial fluids of rheumatoid arthritis patients (Aloe et al. 1992), and in the sera of systemic lupus erythematosus (SLE) patients (Bracci-Laudiero et al. 1993a; Aalto et al. 2002). NGF mRNA expression and protein levels increase in animal models of inflammation such as Freund's adjuvant-induced and carrageenan-induced arthritis (Safieh-Garabedian et al. 1995; Manni and Aloe 1998), in a model of human SLE (Bracci-Laudiero et al. 1996b), and in experimental autoimmune encephalomyelitis (EAE) (De Simone et al. 1996). Many other inflammatory diseases, such as interstitial cystitis (Lowe et al. 1997; Jacobs et al. 2010), allergic asthma (Bonini et al. 1996; Braun et al. 1998), vernal keratoconjunctivitis (Lambiase et al. 1995), Crohn's disease and colitis (di Mola et al. 2000; Stanzel et al. 2008), psoriasis (Fantini et al. 1995; Raychaudhuri et al. 1998), and atopic dermatitis (Toyoda et al. 2002), are characterized by increased tissue and/or circulating NGF. Taken together, these data suggested that inflammation and tissue damage generate mediators that control the local concentration of NGF. Inflammatory cytokines such as IL-1beta, TNF-alpha, and IL-6 are able to modify the basal production of NGF in the organism and induce the synthesis of NGF in a variety of cell types and tissues (Lindholm et al. 1987; Friedman et al. 1990; Gadient et al. 1990; Steiner et al. 1991; Hattori et al. 1994; Kossmann et al. 1996; Marz et al. 1999; Olgart and Frossard 2001; Manni et al. 2003; Moser et al. 2004; Ryan et al. 2008; Seidel et al. 2010). Other inflammatory mediators such as prostaglandins and histamine have also been shown to induce NGF production in certain cell types (Lipnik-Stangelj and Carman-Krzan 2004a, b; Toyomoto et al. 2004; Bullo et al. 2005). In addition, stimulation with specific antigens and cytokines of immune cells involved in innate and acquired immunity modulates their basal expression of NGF (Torcia et al. 1996; Lambiase et al. 1997; Xiang and Nilsson 2000; Caroleo et al. 2001; Kobayashi et al. 2002). Activation of the NF-KB pathway seems to be essential in regulating NGF expression in B lymphocytes (Heese et al. 2006). The endogenous levels of NGF are substantially increased in inflamed tissues, dynamically regulating innervations and neural activities during the inflammatory process (Carr and Undem 2001). During inflammation, the neuronal phenotype can be modified, altering the amount or type of neurotransmitters produced and stored in nerve terminals. As a result, pain sensitivity is increased during inflammation: the properties of the somatosensory neurons change so that the pain threshold decreases, noxious stimuli evoke more acute and prolonged pain (hyperalgesia), and normally innocuous stimuli became painful (allodynia) (Woolf and Salter 2000). This peripheral sensitization increases the firing of nociceptors, and the intense stimulation induces the release from nerve endings of neuropeptides that can influence a variety of immune cell functions. One of the best-known neuroimmune mechanisms involving sensory neuropeptides is neurogenic inflammation: this is characterized by a release of neuropeptides by sensory nerves that in turn induces vasodilatation and plasma extravasation, promotes leukocyte chemotaxis and phagocytosis (Zegarska et al. 2006), and directly affects the release of inflammatory mediators from mast cells and macrophages (Tore and Tuncel 2009). In the mucosae and in lymphoid organs, sensory fibers are in close contact with the cell membranes of mast cells, macrophages, Langerhans cells, and endothelial cells in what has been defined a "neuroimmune junction" (Straub 2004). Neuropeptides released from nerve terminals induce synthesis of inflammatory mediators and cytokines in these cells and affect their functions. While SP has a demonstrated proinflammatory role (O'Connor et al. 2004), CGRP, which is often co-localized with SP in nerve endings, can also have an inhibitory action. CGRP is a potent inhibitor of mitogen- and antigen-stimulated proliferation of T-cells (Carucci et al. 2000), antigen presentation by antigen-presenting cells (Hosoi et al. 1993), and cytokine synthesis (Gomes et al. 2005). The localized increased production of NGF in inflammatory conditions induces modification of gene expression in dorsal root ganglia that can be prevented using anti-NGF antibodies (Donnerer et al. 1992). The neuronal phenotypic changes driven by NGF during inflammation include elevation of neuropeptides, which amplify sensory input signals in the spinal cord and augment neurogenic inflammation in the peripheral tissues.

# 3.2 In Vivo Anti-inflammatory Action

The previously cited evidence indicates that NGF maintains immune cells in a "state of alert" to potential danger signals. NGF by itself does not seem to induce inflammatory and immune responses efficiently, but, in the presence of specific stimuli, it enhances mediator release and cytokine production and activates the functions of innate immune cells. In other words, NGF helps to mount a faster and stronger inflammatory response. However, at present, some of the data obtained in vivo are not consistent with this conclusion. The reasons why NGF production is increased during inflammation and how this enhanced concentration affects the immune and inflammatory responses are still far from being understood. The notion that NGF plays a much more complex role in the scenario of immune response has emerged from studies on experimental models of EAE. In marmoset EAE, intracerebroventricular administration of NGF delays the onset of clinical signs and prevents the full development of histological lesions in the CNS (Villoslada et al. 2000). NGF reduces the number and size of inflammatory infiltrates, and demyelination is minimal in the brain of NGF-treated animals (Villoslada et al. 2000). Moreover, the production of interferon-gamma by the T-cells infiltrating the brain is reduced, while production of IL-10 is enhanced in NGF-treated animals (Villoslada et al. 2000). In a murine model of EAE, intraperitoneal injections of NGF starting one day after immunization with myelin basic



NGF protects against experimental autoimmune encephalomyelitis

**Fig. 4** *NGF protects common marmosets against autoimmune encephalomyelitis.* The figure is a summary of results published by Villoslada and collaborators (Villoslada et al. 2000). In this paper, the authors demonstrated that intracerebroventricular (icv) NGF was able to revert clinical manifestations of the experimental autoimmune encephalomyelitis (EAE), an animal model for human multiple sclerosis. In such a model, the immunization with myelin oligodendrocyte glycoprotein (MOG) induces a massive autoimmune response in the central nervous system, characterized by infiltration of activated T-cells and activation of glial cells. Brain-delivered NGF was able to delay the onset of clinical signs and prevent the full development of histological lesions, reducing the number and size of inflammatory infiltrates and the demyelination. Moreover, the production of interferon-gamma by the T-cells infiltrating the brain was reduced, while production of IL-10 by glial cells was enhanced in NGF-treated animals

protein (MBP) or after adoptive transfer of encephalitogenic T-cells led to a delayed onset, decreased clinical signs of the disease, and enhanced survival of the animals (Arredondo et al. 2001). In a T-cell transfer model of EAE, myelin basic protein-specific CD4+ T-cell clones were transfected with recombinant retrovirus encoding for NGF and used to induce EAE in rats (Flugel et al. 2001). The clones producing NGF did not induce EAE, while the pathogenic wild-type clones injected together with the NGF-transfected clones induced a very mild disease with no or only minor clinical signs (Flugel et al. 2001). A reduction in inflammatory cells crossing the endothelial blood-brain barrier (BBB), with the number of activated macrophages being especially decreased, was also found (Flugel et al. 2001). Complementary effects are observed in another EAE rat model, in which NGF deprivation results in exacerbated brain inflammation and more severe clinical signs (Micera et al. 2000). Thus, the overall data obtained from EAE models point to an anti-inflammatory role of NGF in CNS immune-mediated inflammation. NGF seems to exert its effects directly on immune cell activity: inhibition of monocyte migration and antigen presentation, altered T-helper balance, downregulation of IFN-gamma synthesis, and upregulation of the antiinflammatory cytokine IL-10 (Fig. 4). Similarly, in experimental models of colitis, it has been shown that NGF production remains elevated in the tissue until colon inflammation subsides (Barada et al. 2007), while neutralizing NGF by anti-NGF antibodies causes a significant increase in the severity of clinical signs with more

extended lesions and ulcers, together with an increase in the number of infiltrating neutrophils and macrophages (Reinshagen et al. 2000). Also in this experimental model, the increase in NGF correlates with the enhancement of IL-10 (Barada et al. 2007), possibly because NGF and IL-10 are able to regulate each other's expression in intestinal epithelial cells (Ma et al. 2003). In addition, functional sensory neurons and their neuropeptides (i.e., CGRP) are probably involved in the in vivo action of NGF. In the colitis model, enhanced tissue inflammation caused by NGF neutralization is associated with a marked reduction in CGRP content in the gut (Reinshagen et al. 1994). Moreover, the ablation of sensory fibers results in a marked severity of inflammation in acute and chronic models of experimental colitis, and in both cases the decrease in CGRP content or its inhibition in the colon using CGRP receptor antagonist aggravates experimental colitis (Reinshagen et al. 1998).

The NGF produced in vivo at the site of inflammation thus represents a critical link between nervous and immune cells. This hypothesis is supported by studies on a mouse model in which the animal's skin received ultraviolet B (UVB) irradiation. UVB induces systemic suppression of contact hypersensitivity (CHS) responses, and NGF, together with nerves and neuropeptides, seems to play a relevant role in this phenomenon. In mouse epidermis, UVB irradiation upregulates NGF expression, which peaks after 12 h (Gillardon et al. 1995). Anti-NGF antibodies administered prior to UVB irradiation abrogate the systemic suppression of CHS in mice (Townley et al. 2002). The CHS response in healthy mice was also inhibited by NGF treatment of the skin prior to trinitrochlorobenzene sensitization (Townley et al. 2002). Moreover, when mice were treated with capsaicin, which depletes nerve endings of sensory neuropeptides, preincubation with NGF was not effective in suppressing CHS response (Townley et al. 2002). Also in this model, NGF-induced synthesis of CGRP seems to represent a key mediator for the control of immune cell activity and for balancing pro- and anti-inflammatory responses. In UV-induced tolerance, the induction of CGRP synthesis in skin after UV radiation (Legat et al. 2004) is regulated by keratinocyte-derived NGF (Townley et al. 2002). In vivo CGRP administration dampens the immune response essentially by affecting antigen presentation in a variety of antigen-presenting cells (Raud et al. 1991; Hosoi et al. 1993; Asahina et al. 1995; Carucci et al. 2000), through inhibition of the NF-kappaB pathway (Millet et al. 2000) and a rapid upregulation of inducible cAMP early repressor (Harzenetter et al. 2007), which causes premature repression of inflammation-induced transcriptional activity. Thus, the ability of NGF in vivo to regulate production and release of sensory neuropeptides seems to be a key factor not only in keeping the defense system active (i.e., through SP) but also in activating inhibitory pathways that limit inflammation.

Data from in vivo models of inflammation show that integrity of the peripheral nervous system is essential for the activation of an anti-inflammatory feedback circuit (Rook et al. 2002; Holzer 2007; Dirmeier et al. 2008; Boisse et al. 2009; Kroeger et al. 2009). Cutaneous infections and altered immune responses in diabetes represent a characteristic example of nerve damage leading to uncontrolled

tissue inflammation. Impaired production of NGF is responsible for the reduction in skin innervations in diabetes (Hellweg and Hartung 1990; Anand 1996). In animal models, the exogenous administration of NGF, while improving nerve functions and restoring innervation in the skin (Christianson et al. 2003), normalizes the diabetes-impaired response to wound healing by decreasing neutrophil accumulation and increasing reepithelialization and matrix density (Matsuda et al. 1998).

#### 4 Conclusion

Since its discovery in the 1950s, NGF has been extensively studied for its trophic, pro-survival, and differentiative effects in both neuronal and nonneuronal cell populations. Several studies have shown that the neurotrophin can display contradictory behavior, acting either as a survival or as an apoptotic factor in neuronal cells (Lee et al. 2001). In the last decade, the apparently controversial effects of NGF on neuronal cells have been better elucidated, in light of the discovery and characterization of the precursor form of NGF, named proNGF. ProNGF can be released by cells and cleaved to form mature NGF (Bruno and Cuello 2006) or can act as an active molecule, being a specific and selective ligand for a receptor complex made by p75<sup>NTR</sup> and sortilin (Fahnestock et al. 2004), a protein that shuttles other proteins within cells and acts as a receptor for neurotensin (Mazella 2001). In several injury models in the central nervous system, proNGF expression is detectable for several days to weeks following injury. Surprisingly, little conversion of proNGF to mature NGF is observed in these in vivo settings (Beattie et al. 2002; Harrington et al. 2004; Jansen et al. 2007). This finding suggests that the mature and pro-forms of NGF may perform opposing actions. Expression of the  $p75^{NTR}$  receptor has emerged as a key regulatory element in proNGF-induced cell death. In most adult tissues, p75<sup>NTR</sup> is expressed at low levels, in contrast to its stronger and more widespread distribution in development (Yang et al. 2009; Roux and Barker 2002). However, in pathologic states, including seizure, brain injury, ischemia, and excitotoxicity, p75<sup>NTR</sup> expression is induced. The significant reduction in injury-induced apoptosis observed in p75<sup>NTR</sup>-deficient mice (Troy et al. 2002; Harrington et al. 2004) clearly indicates the importance of p75<sup>NTR</sup> induction in determining cell loss following injury.

Until now, very few studies have addressed the function of proNGF in the immune system, but this promises to be an exciting field of research for the near future. Only very recently was the action of proNGF on immune cells addressed in a work investigating the effects of  $p75^{NTR}$ /sortilin activation in NK cells in vitro (Rogers et al. 2010). Freshly isolated NK and B-cells, but not T lymphocytes, expressed significant amounts of sortilin but did not express significant amounts of  $p75^{NTR}$  in unstimulated conditions. The expression of  $p75^{NTR}$  is induced in NK cells by IL-12. Addition of proNGF in the presence of IL-12 induced NK cell death (Rogers et al. 2010). Immune cells, such as mast cells, have been indicated as a possible source of proNGF. Indeed, proNGF

immunoreactivity was increased in skin-homed mast cells in an in vivo model of neuropathic pain (Peleshok and Ribeiro-da-Silva 2012). Mast cells may also have a role as key players in the extracellular maturation of NGF from its precursor. Tryptase-secreting mast cells have been detected in close proximity to NGF-secreting cells in the testis of infertile patients, and it is likely that they are able to alter the ratios of proNGF/NGF by increasing tryptase enzymatic cleavage of proNGF (Spinnler et al. 2011). There is also evidence that neuroimmune modulators may be involved in regulating the proNGF/NGF balance. In an in vitro model of inflamed skin, the addition of neuropeptides such as SP, CGRP, VIP, and GAL to normal human keratinocytes results in an increased expression of IL-1 $\beta$ , IL-8, and TNF- $\alpha$ ; release of proNGF; and its increased conversion into mature NGF in the culture medium (Dallos et al. 2006).

In view of the apparent discrepancies in the effects of NGF on immune functions, it is to be hoped that future studies of NGF and proNGF will help to clarify the role of this neurotrophin in regulating immune response and inflammation. A better understanding of proNGF/NGF actions could be of immense value, with translational implications for human diseases.

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# NGF/P75 in Cell Cycle and Tetraploidy

Noelia López-Sánchez, María Carmen Ovejero-Benito, Claudia Rodríguez-Ruiz, and José María Frade

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### Abstract

Neurotoxicity, neurodegeneration, and other disorders affecting neuron survival are often related to cell cycle reentry in neurons. Traditionally, cell cycle reentry of these postmitotic cells has been thought to be associated with apoptosis. Nevertheless, cell cycle reentry and DNA synthesis in neurons could also lead to tetraploidy which may explain long-lasting neurodegenerative processes. During development, a subpopulation of newborn neurons reactivates the cell cycle and becomes tetraploid in response to p75<sup>NTR</sup> activation. These neurons enlarge their cell bodies and increase their dendritic trees, thus generating specific neuronal populations that innervate particular areas. Pathological states in the nervous system could also lead to p75<sup>NTR</sup>-dependent neuronal tetraploidy.

N. López-Sánchez • M.C. Ovejero-Benito • C. Rodríguez-Ruiz • J.M. Frade (⊠) Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC), Madrid, Spain e-mail: nlopez@cajal.csic.es; mcovejerob@cajal.csic.es; claudia.roru@gmail.com; frade@cajal.csic.es

De novo tetraploid neurons might become structurally and functionally altered, thus leading to neurodegeneration at late stages of the disease. This chapter describes what is currently known about the interplay between  $p75^{NTR}$  and the cell cycle, stressing the role played by different  $p75^{NTR}$  interactors, including the MAGE and Bex1/NADE adaptor proteins and the transcription factors SC1, NRIF, and Sall2, in cell cycle regulation. The chapter also discusses on the effects of  $p75^{NTR}$ , as a cell cycle regulator, in neural stem cell proliferation, neurogenesis, and neuronal tetraploidization, as well as on the connection of  $p75^{NTR}$  in pathology, including its putative effects in neurodegeneration.

List of Abb	reviations					
$A\beta$ Amyloid- $\beta$						
BDNF	Brain-derived neurotrophic factor					
Bex-1	Brain-expressed X-linked 1					
BrdU	Bromodeoxyuridine					
cdk	Cyclin-dependent kinase					
CMAGE	Chicken MAGE					
CNS	Central nervous system					
Dlxin-1	Dlx/Msx-interacting MAGE/Necdin family protein					
E2F1	E2 promoter-binding factor-1					
EGFP	Enhanced green fluorescent protein					
ERK	Extracellular signal-regulated kinase					
JNK	c-Jun N-terminal kinase					
MAGE	Melanoma antigen					
MAPK	Mitogen-activated protein kinase					
NADE	p75 <sup>NTR</sup> -associated cell death executor					
NGF	Nerve growth factor					
NRAGE	Neurotrophin receptor-interacting MAGE homolog					
NRIF	Neurotrophin receptor-interacting factor					
NT3	Neurotrophin-3					
p75 <sup>ICD</sup>	p75 <sup>NTR</sup> intracellular domain					
p75 <sup>NTR</sup>	p75 neurotrophin receptor					
PCNA	Proliferating cell nuclear marker					
PNS	Peripheral nervous system					
PR/SET	Positive regulatory/suppressor of variegation, enhancer of zeste,					
	trithorax					
Rb	Retinoblastoma					
RGCs	Retinal ganglion cells					
Sall2	Sal-like protein 2					
SC1	Schwann cell factor 1					
SGZ	Subgranular zone					
SVZ	Subventricular zone					
TNF	Tumor necrosis factor					
Trk	Tropomyosin-related kinase					

### 1 Introduction

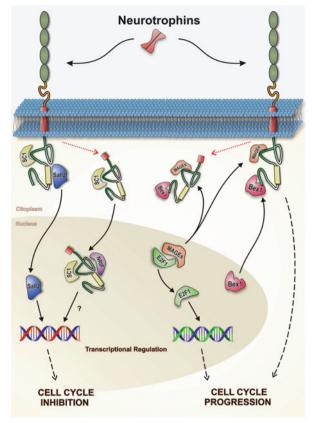
Neurotrophins are best known as neuronal survival factors with multiple effects in the structure and function of the adult vertebrate nervous system (Huang and Reichardt 2001). The first neurotrophin to be described was NGF in the early 1950s (Levi-Montalcini and Hamburger 1951), being initially purified from sarcoma tumor cells (Cohen et al. 1954). Several decades later, other members of the neurotrophin family were characterized, including BDNF (Barde et al. 1982), NT3 (Hohn et al. 1990; Maisonpierre et al. 1990; Rosenthal et al. 1990; Jones and Reichardt 1990), and NT4/5 (Hallböök et al. 1991; Berkemeier et al. 1991). In the brain, mature neurotrophins coexist with their precursor pro-forms, namely, proNGF, proBDNF, and proNT3 (Fahnestock et al. 2001; Michalski and Fahnestock 2003). The neurotrophin pro-forms fulfill specific functions, usually opposed to that of the mature neurotrophins, including induction of cell death (Lee et al. 2001; Teng et al. 2005; Yano et al. 2009) and, in the case of proBDNF, synapse modulation (Lu 2003; Woo et al. 2005; Greenberg et al. 2009). These opposed effects can be explained due to the capacity of neurotrophins to interact with two different types of receptors, p75<sup>NTR</sup> and the members of the Trk tyrosine kinase receptor family. p75<sup>NTR</sup> was the first member of the TNF receptor family to be described (Chao et al. 1986; Johnson et al. 1986; Radeke et al. 1987). Unlike the Trk receptors, known to interact with specific neurotrophins (Kaplan et al. 1991; Klein et al. 1991a, b, 1992; Lamballe et al. 1991), p75<sup>NTR</sup> can bind to, and be activated by, all neurotrophins (Rodríguez-Tébar et al. 1990, 1992). Although p75<sup>NTR</sup> was initially thought to be a mere Trk co-receptor, it is now clear that this receptor is at the core of at least two other signaling networks, including the p75<sup>NTR</sup>/sortilin platform that induces apoptosis in response to the pro-forms of the neurotrophins (Nykjaer et al. 2004; Teng et al. 2005; Yano et al. 2009) and the p75<sup>NTR</sup>/NgR/LINGO-1 complex that regulates axonal outgrowth (Wang et al. 2002; Mi et al. 2004). In some cases, signaling through p75<sup>NTR</sup> requires the release of its intracellular domain (p75<sup>ICD</sup>), which can be translocated to the nucleus (Frade 2005; Parkhurst et al. 2010). This mechanism has been shown to lead to apoptosis (Majdan et al. 1997; Kenchappa et al. 2006, 2010; Podlesniy et al. 2006), and it can also participate in Trk signaling (Skeldal et al. 2011).

Mounting evidence indicates that neurotrophins can regulate cell cycle progression through both  $p75^{NTR}$  and the Trk receptors (López-Sánchez and Frade 2002). This chapter will firstly describe what is known about the interplay between  $p75^{NTR}$  and the cell cycle machinery. Then, the chapter will be focused on the role played by this receptor in the normal nervous system in terms of cell cycle regulation (proliferation of neural stem cells, embryonic and adult neurogenesis, and neuronal tetraploidization). Finally, a discussion on the possible involvement of  $p75^{NTR}$  as a cell cycle regulator in neurodegeneration will be presented.

# 2 Interplay Between p75<sup>NTR</sup> and the Cell Cycle

 $p75^{NTR}$  is the founder member of the tumor necrosis factor receptor family. This receptor, characterized by the presence of one transmembrane domain and a type II death domain within its intracellular domain, is expressed by a variety of neuronal and non-neuronal cell types, frequently showing proliferative capacity. In this regard, p75<sup>NTR</sup> was initially described in proliferating cells derived from pheochromocytoma, neuroblastoma, and melanoma tumors (Yankner and Shooter 1982). More recently, this receptor is often used as a marker of neural progenitor cells in the adult brain (Giuliani et al. 2004; Young et al. 2007; Bernabeu and Longo 2010) and as a tumor suppressor gene in a number of tumor cells (Krygier and Djakiew 2001; Yuanlong et al. 2008; Jin et al. 2007; Khwaja and Djakiew 2003; Khwaja et al. 2006). p75<sup>NTR</sup> has opposing effects on the cell cycle, inducing cell cycle arrest in some instances and cell cycle progression in other circumstances. For instance, p75<sup>NTR</sup>-dependent cell cycle arrest, associated with reduced expression of cyclin D1, cyclin E, cdk2, E2F1, PCNA, and the cdk inhibitor p16<sup>INK4a</sup>, decreased cdk2 activity, and hyper-phosphorylation of Rb protein, has been shown to occur in bladder tumor epithelial cells (Khwaja and Djakiew 2003). In human gastric cancer cells, p75<sup>NTR</sup> expression suppresses proliferation by downregulation of cyclin A, cyclin D1, cyclin E, cdk2, and phospho-Rb and upregulation of the cell cycle inhibitors Rb and p27<sup>KIP1</sup>. This antiproliferative activity in human gastric cancer cells is dependent on the presence of the death domain of  $p75^{NTR}$  (Jin et al. 2007). p75<sup>NTR</sup> expression in hepatocellular carcinoma cells has also been shown to downregulate the expression of cyclin A, cyclin D1, cyclin E, cdk2, p-Rb, and PCNA and upregulate the expression of the cell cycle inhibitor Rb (Yuanlong et al. 2008). Examples of p75<sup>NTR</sup>-dependent cell cycle progression are also available in the literature. For instance,  $p75^{NTR}$  has been shown to be expressed by human oral keratinocyte stem cells showing high in vitro proliferative capacity and clonal growth potential (Nakamura et al. 2007) as well as by human esophageal keratinocyte stem cells able to undergo self-renewal and self-amplification (Okumura et al. 2003). Moreover, p75<sup>NTR</sup> can be detected in cycling cells from the subventricular zone of adult male rats (Giuliani et al. 2004), and the addition of NGF has been shown to revert p75<sup>NTR</sup>-dependent cell cycle arrest in bladder tumor epithelial cells (Khwaja and Djakiew 2003).

The mechanisms used by  $p75^{NTR}$  to regulate cell cycle progression are beginning to be elucidated. In this regard,  $p75^{NTR}$  is known to activate different members of the MAPK family able to regulate the cell cycle (Ambrosino and Nebreda 2001; Zhang and Liu 2002), including JNK,  $p38^{MAPK}$ , and ERK (Casaccia-Bonnefil et al. 1996; Susen et al. 1999; Costantini et al. 2005; Jiang et al. 2007). Moreover, the opposing effects of  $p75^{NTR}$  on cell cycle regulation can be explained by the different molecules with which this receptor interacts (see Fig. 1).  $p75^{NTR}$  can modulate the cell cycle through the interaction of its intracellular domain with a number of proteins known to regulate the cell cycle, some of which are transcription factors. In some instances,  $p75^{NTR}$ has been shown to sequester these molecules at the plasma membrane. Fig. 1 Scheme of the signaling pathways used by  $p75^{NTR}$  to regulate the cell cycle. Dotted arrows represent the release of  $p75^{ICD}$  as a consequence of regulated y-secretasedependent cleavage of p75<sup>NTR</sup>. Solid arrows describe the dynamics of the different proteins involved in the p75<sup>NTR</sup> signaling. Dashed arrows represent the final biological output of the corresponding pathways. The *auestion mark* represents the putative transcriptional activity of p75<sup>ICD</sup> complexed with NRIF and/or SC1. See main text for further details



In addition, p75<sup>NTR</sup> is susceptible to undergo  $\gamma$ -secretase-dependent cleavage and release its intracellular domain (p75<sup>ICD</sup>) which may subsequently translocate to the nucleus (Kanning et al. 2003; Frade 2005). The interaction of p75<sup>ICD</sup> with transcription factors led to the suggestion that it can regulate gene expression when located in the nucleus. This notion has recently been confirmed as p75<sup>ICD</sup> has been shown to translocate to the nucleus and activate a green fluorescent protein reporter (Parkhurst et al. 2010). Furthermore, NGF can induce the interaction of p75<sup>ICD</sup> with the cyclin E promoter, thus inhibiting its activity in HeLa cells (Parkhurst et al. 2010), a result consistent with the observation that NGF-dependent nuclear translocation of p75<sup>ICD</sup> in schwannoma cells (Frade 2005) correlates with cell cycle repression in these cells (Morillo and Frade 2008). The expression of p75<sup>ICD</sup> in p75<sup>NTR</sup>-negative PC12 cells downregulates cyclin D2 expression as well (Fritz et al. 2006). p75<sup>NTR</sup> can regulate the cell cycle by means of recruitment of adaptor proteins such as different MAGE proteins or Bex1, known to be involved in the cell cycle regulation, and also by regulating the expression of cell cycleregulating genes through transcription factors such as SC1, NRIF, and Sall2, all of them able to interact with  $p75^{ICD}$ .

### 2.1 MAGE Proteins

The MAGE protein family comprises more than 60 adaptor proteins in mammals, containing a conserved MAGE homology domain. Originally, MAGE proteins were discovered because peptides derived from MAGE gene products are presented on the cell surface of human melanoma cells by the major histocompatibility complex (Barker and Salehi 2002). MAGE proteins can be divided into two major groups: type I and type II MAGE proteins. The former are only expressed in tumors, male germinal cells, and placenta. In contrast, the latter group, defined by a phylogenetically distinct MAGE homology domain, comprises around ten different proteins expressed in both differentiating and adult cells (Barker and Salehi 2002). Most of the MAGE proteins are encoded by genes containing a single exon. suggesting that the whole MAGE protein family evolved from a retrotransposition event that occurred in an ancestral gene orthologue of the gene encoding NRAGE/ Dlxin-1/MAGE-D1, the only MAGE gene containing exons in its sequence, followed by extensive gene duplication. This hypothesis is consistent with the existence of a single MAGE gene in all nonmammalian species including unicellular, plants, fungi, invertebrates, and nonmammalian vertebrates (López-Sánchez et al. 2007).

The mammalian MAGE II proteins Necdin, NRAGE/Dlxin-1/MAGE-D1, MAGE-H1, and MAGE-G1 as well as CMAGE are known to interact with p75<sup>NTR</sup> through its intracellular domain (Salehi et al. 2000; Tcherpakov et al. 2002; Kuwako et al. 2004; López-Sánchez et al. 2007), and they have effects on cell cycle regulation. Therefore, these proteins are candidates for the effects of  $p75^{NTR}$  on the cell cycle. Necdin can bind to the transcription factor E2F1, thus blocking its capacity to trigger G1/S transition (Taniura et al. 1998; Kuwako et al. 2004). The activation of p75<sup>NTR</sup> by NGF can recruit Necdin to the cell membrane, thus favoring E2F1-dependent cell cycle progression and subsequent death of differentiated neurons (Kuwako et al. 2004) (see Fig. 1). Alternatively, the presence of the p75<sup>ICD</sup> fragment can displace both Necdin and CMAGE from their interaction with E2F1 in differentiating neurons, thus leading to cell E2F1-dependent cycle reentry and apoptosis (López-Sánchez et al. 2007) (see Fig. 1). NRAGE/Dlxin-1/MAGE-D1 can also block cell cycle progression (Salehi et al. 2000), likely by inducing the expression of the cell cycle-inhibiting protein p21<sup>Waf1</sup> in a p53-dependent manner (Wen et al. 2004). NRAGE/Dlxin-1/ MAGE-D1 associates with the plasma membrane when NGF is bound to p75<sup>NTR</sup> (Salehi et al. 2000), as it occurs with Necdin (see above). Therefore, sequestering of MAGE proteins by p75<sup>NTR</sup> seems to be a general mechanism for enhancement of cell cycle progression in different cellular systems.

### 2.2 Bex/NADE

p75<sup>NTR</sup> is also known to interact through its death domain with the small adaptor-like proteins from the Bex/NADE family (Mukai et al. 2000; Vilar et al. 2006), including Bex1 and Bex3/NADE. Among them, Bex1 can constitutively

interact with p75<sup>NTR</sup> (see Fig. 1) and favor cell cycle progression when overexpressed in differentiating PC12 cells (Vilar et al. 2006). This is consistent with the increase of Bex1 during S phase in PC12 cells, as it occurs with the presence of  $p75^{NTR}$  on the surface of these same cells (Urdiales et al. 1998). Upon NGF treatment, Bex1 translocates from the nucleus to the cytoplasm (see Fig. 1), suggesting that this mechanism is required for the effect of this protein on cell cycle regulation. Interestingly, in other cell systems Bex1/2 seems to act as a putative tumor suppressor gene in malignant glioma (Foltz et al. 2006), although in this case the participation of  $p75^{NTR}$  is currently unknown.

### 2.3 SC1

The p75<sup>NTR</sup>-interacting protein SC1 is a member of the PR/SET domain-containing zinc finger protein family, which can be detected in both the nucleus and cytoplasm when overexpressed (Chittka and Chao 1999). SC1 is redistributed from the cytoplasm to the nucleus after NGF treatment in a p75<sup>NTR</sup>-dependent manner, acting as a transcriptional repressor (see Fig. 1). Nuclear SC1 correlates with the loss of BrdU incorporation, likely due to its ability to repress cyclin E expression (Chittka et al. 2004). Interestingly, NGF has been shown to induce the association of endogenous p75<sup>ICD</sup> with the cyclin E promoter (Parkhurst et al. 2010), suggesting that a complex formed by p75<sup>ICD</sup> and SC1 can be formed in the promoter of different genes encoding cell cycle-regulating proteins (see Fig. 1).

### 2.4 Sall2

Sall2 is another transcription factor able to regulate cell cycle progression which can interact with  $p75^{NTR}$  through its intracellular domain (see Fig. 1). This protein is a tumor suppressor that becomes dissociated from its binding to  $p75^{NTR}$  upon interaction of the latter with NGF (Pincheira et al. 2009). Sall2 is known to activate  $p21^{Waf1}$  and promote growth arrest in neurons. This observation is consistent with the  $p75^{NTR}$ -dependent increase of  $p21^{Waf1}$  expression and decrease of Rb phosphorylation in breast cancer cells (Verbeke et al. 2010).

### 2.5 NRIF

The zinc finger proteins of the Krüppel family NRIF1 and NRIF2 are able to interact with the intracellular domain of  $p75^{NTR}$  through its juxtamembrane and death domains (Casademunt et al. 1999; Benzel et al. 2001) (see Fig. 1). These proteins have been suggested to be transcription factors based on structural characteristics and their ability to translocate to the nucleus (Gentry et al. 2004), and they prevent cell cycle progression when expressed in 293T cells. Interestingly, the release of  $p75^{ICD}$  correlates with the translocation of NRIF to the nucleus

(Kenchappa et al. 2006; Volosin et al. 2008), suggesting the complex NRIF/ $p75^{ICD}$  may participate in the regulation of the cell cycle, as it occurs with  $p75^{ICD}$  and SC1 (see above).

# 3 The Role of p75<sup>NTR</sup> as a Cell Cycle Regulator in Nervous System Physiology

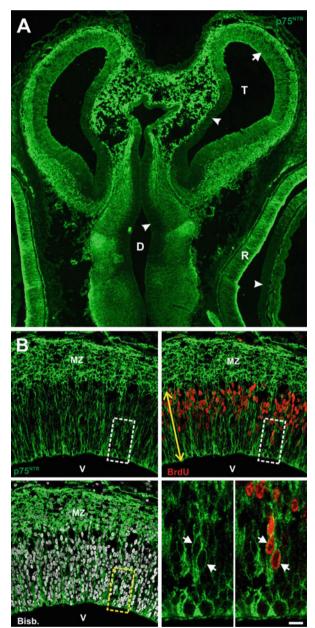
A critical feature of the nervous system is the requirement of neurons to be postmitotic. The avoidance of cell division in these cells is likely derived from their complex morphology, required for their function, which is incompatible with the reorganization of the cytoskeleton necessary for the mitotic process. Therefore, at certain point of neural development, proliferating neuronal precursors stop dividing and differentiate as neurons in a process referred to as neurogenesis. During development, p75<sup>NTR</sup> can be detected in the proliferating epithelium of the retina (Frade and Barde 1999; Morillo et al. 2010; López-Sánchez et al. 2011), spinal cord (Frade and Barde 1999; López-Sánchez et al. 2011), and other structures of the CNS including the dorsal telencephalon in the chick (Fig. 2), suggesting that it can participate in the neurogenic process. Furthermore,  $p75^{NTR}$  is expressed by neural crest cells (Bannerman and Pleasure 1993), a proliferating cell population giving rise to the PNS, among other structures. Some evidence exists for the participation of p75<sup>NTR</sup> in cell cycle withdrawal of precursor cells and neuronal differentiation during embryonic development (Hapner et al. 1998; Hosomi et al. 2003; Zhang et al. 2009). Much more evidence exists for the participation of p75<sup>NTR</sup> in adult neurogenesis and neuronal tetraploidization during embryonic development. These two processes are further described below.

### 3.1 Adult Neurogenesis

For many decades it was assumed that the generation of functional neurons from precursors (i.e., neurogenesis) only occurs during embryonic and perinatal stages of development in mammals. This view has been ruled out, and nowadays it is widely accepted that neural stem cells can be present in the adult nervous system, being able to proliferate in vitro giving rise to neurospheres (Reynolds and Weiss 1992). Neurons are known to be produced during adulthood in two main brain regions, in the SVZ of the lateral ventricle (giving rise to the rostral migratory stream that colonizes the olfactory bulb) and the SGZ from the dentate gyrus of the hippocampus (producing dentate granule neurons).

Interestingly, p75<sup>NTR</sup> is expressed by cycling cells in the SVZ of adult male rats (Giuliani et al. 2004). This p75<sup>NTR</sup>-positive cell population integrates all of the neurosphere-producing SVZ precursors, which are known to enhance neuronal production in response to NGF or BDNF (Young et al. 2007). The importance of p75<sup>NTR</sup> in SVZ adult neurogenesis is further supported by the observation that the neurogenic potential is reduced in neurospheres derived from p75<sup>NTR</sup> knockout

**Fig. 2**  $p75^{NTR}$  expression in the chick brain at developmental day 6 (E6). (A) Horizontal section through the E6 chick brain, immunostained with the ChEX anti-p75<sup>NTR</sup> antibody (kindly provided by L. F. Reichardt, University of California San Francisco). p75<sup>NTR</sup> is expressed in specific regions of the neuroepithelium (arrow), while this protein cannot be detected in other areas of the neuroepithelium (arrowheads). T telencephalic derivatives, D diencephalic derivatives, *R* retina. (**B**) A high magnification from the telencephalic wall (dorsal hyperpallium) from E6 chick embryos treated with BrdU for 20 min. BrdU is incorporated by nuclei of progenitors undergoing S phase (red) which, due to the interkinetic nuclear movement, are mostly located far from the ventricle (V). Nevertheless, some nuclei from p75<sup>NTR</sup>-expressing precursors incorporate BrdU when located close to the ventricle (dashed box and arrows in the magnification of the dashed box shown in the bottom right panels). Nuclei were labeled with bisbenzimide (Bisb.). Abbreviation: MZ mantle zone. Bar  $-65 \mu m (A)$ , 25 µm (**B**)



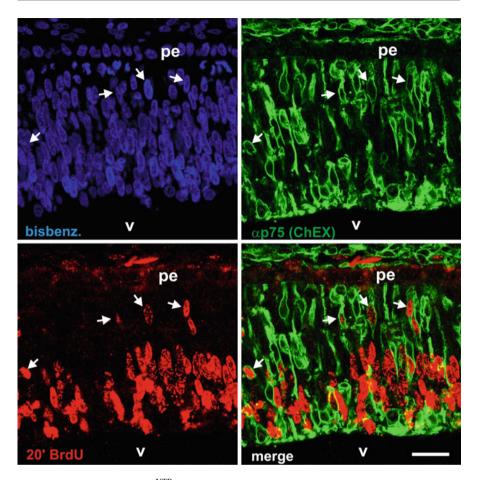
mice (Young et al. 2007).  $p75^{NTR}$  might also regulate proliferation of SVZ precursors in pathological states. For instance,  $p75^{NTR}$  activation with  $A\beta_{1-42}$ , the pathological form of  $A\beta$  in Alzheimer's disease, has been shown to favor proliferation and neuronal differentiation in neurosphere-forming adult progenitors,

an effect that is not observed in p75<sup>NTR</sup> knockout mice (Sotthibundhu et al. 2009). Moreover, an increase in the proliferation activity of p75<sup>NTR</sup>-positive SVZ precursors has been described in rats subjected to experimental allergic encephalomyelitis, a demyelinating disease widely used as an experimental model for human multiple sclerosis (Calzà et al. 1998). In the SGZ, p75<sup>NTR</sup> is expressed in dividing SGZ cells (Okano et al. 1996; Bernabeu and Longo 2010), and it seems to participate in adult hippocampal neurogenesis as neurogenic potential is reduced in the SGZ of p75<sup>NTR</sup> knockout mice (Bernabeu and Longo 2010; Colditz et al. 2010), maybe due to reduced survival of neuroblasts (Catts et al. 2008). Neurogenesis can also take place in different regions of the nervous system in response to injury. This is the case of the adult dorsal root ganglia, in which a subpopulation of glia may play a role in neurogenesis after peripheral nerve injury. These cells express p75<sup>NTR</sup>, and in vitro they have been reported to form neurospheres (Li et al. 2007). Neurogenesis in the striatum in response to focal cerebral ischemia can be also potentiated by p75<sup>NTR</sup> since the proportion of neurons incorporating BrdU was increased in this paradigm (Zhu et al. 2011).

### 3.2 Neuronal Tetraploidy

The acquisition of multicellularity in evolution has allowed the emergence of the somatic lineage, in which genomic modifications may occur without transmission to the descendants. These genomic alterations, including somatic polyploidy, may participate in the differentiation of specific tissues. This is the case, for instance, of the liver and heart in vertebrates, which have both been shown to contain polyploid cells (Anatskaya and Vinogradov 2007).

Examples of somatic polyploidy in neurons have been described in some invertebrates (Coggeshall et al. 1970: Manfredi Romanini et al. 1973). In contrast, the presence of polyploid neurons in the vertebrate nervous system has been under controversy for decades (see Swartz and Bhatnagar 1981). It was not until the advent of modern techniques such as fluorescent in situ hybridization, flow cytometry, slide-based cytometry, and quantitative PCR analysis of nuclear DNA that the classical belief that all vertebrate neurons contain a diploid DNA amount (Swift 1953) was challenged. Recent analyses, for instance, have proved that 10 % of cortical neurons display a more than diploid DNA content in humans (Mosch et al. 2007). In this regard, tetraploid neurons have been shown to exist in different neural structures (López-Sánchez et al. 2011) including the normal vertebrate retina. In this latter tissue, tetraploid neurons constitute a subpopulation of RGCs that, in the chick, innervate lamina F in the stratumgriseum-et-fibrosum-superficiale of the tectal cortex (Morillo et al. 2010). In the retina, tetraploid RGCs are generated as they migrate to the ganglion cell layer, soon after they undergo their final mitosis and acquire neuronal markers (Morillo et al. 2010). During this stage differentiating RGCs express p75<sup>NTR</sup> and a subpopulation of these neurons incorporates BrdU, thus becoming tetraploid



**Fig. 3** Colocalization of  $p75^{NTR}$  and BrdU in apically located cells from the E6 chick retina. Embryos were treated for 15 min with 40 µl of a solution containing 10 µg/ml BrdU and then fixed. Sagittal sections through the E6 chick retina were obtained, and they were subsequently immunostained with the ChEX anti- $p75^{NTR}$  antibody (*green*) and an anti-BrdU antibody (*red*) as previously described (López-Sánchez et al. 2011).  $p75^{NTR}$  is expressed by BrdU-positive cells with nuclei located at the apical portion of the neuroepithelium (*arrows*). Nuclei were labeled with bisbenzimide (*Bisb.*). Abbreviations: *pe* pigment epithelium, *v* vitreous body. Bar, 18 µm

throughout an endoreduplicative mechanism (Morillo et al. 2010; see also Fig. 3). Our work has demonstrated that DNA duplication in differentiating RGCs is triggered by NGF through p75<sup>NTR</sup>, since blocking antibodies against these molecules were able to prevent cell cycle reentry (Morillo et al. 2010). BrdU incorporation in neural progenitor cells expressing p75<sup>NTR</sup> can also be observed in other neural structures of the chick embryo such as the spinal cord (López-Sánchez et al. 2011) or the telencephalon (Fig. 2b). Although further analyses are still needed to demonstrate that these p75<sup>NTR</sup>-positive cells are equivalent to those neurons

becoming tetraploid in the retina (Morillo et al. 2010), these results suggest that the participation of  $p75^{NTR}$  in neuronal tetraploidization is a general feature of the developing vertebrate nervous system.

Somatic polyploidy can be generated by different mechanisms usually linked to alterations of the mitotic cycle, including endoreduplication (i.e., a modified version of the cell cycle characterized by S phase without mitosis) and endomitosis (i.e., a mitotic cycle in the absence of anaphase/cytokinesis thus resulting in genomic DNA duplication) (Edgar and Orr-Weaver 2001; Ullah et al. 2009). Tetraploid RGCs are originated in the developing retina through an endoreduplicative cycle since they remain in a G2-like state after replicating their nuclear DNA (Morillo et al. 2010). Evidence exists that the neurotrophin BDNF, through its receptor TrkB, fulfills a critical role in preventing G2/M transition in retinal tetraploid neurons. Indeed, blockade of endogenous BDNF resulted in an increase of differentiating RGCs undergoing mitosis (Morillo et al. 2010). Therefore, neurotrophins, including NGF and BDNF, seem to play a critical role in neuronal tetraploidization in vertebrates. Tetraploid neurons that undergo mitosis finally die (Frade 2000; Morillo et al. 2010; López-Sánchez et al. 2011), and the regulation of the G2/M transition in these cells is likely crucial for removal of specific neuronal types during development (Frade et al. 1997), thus adjusting the ratio between tetraploid and diploid neurons in specific areas. Importantly, this process may be involved in neuronal death associated to neurodegenerative diseases (see below).

Polyploidy is usually linked to enlarged cell size (Edgar and Orr-Weaver 2001; Ullah et al. 2009), a concept that can be extrapolated to the vertebrate nervous system. For instance, neuronal hypertrophy has been shown to occur in tetraploid strains of Xenopus laevis, whose neurons contain significantly enlarged cell somas and dendrites (Szaro and Tompkins 1987). As expected, our studies have revealed that tetraploid RGCs show enlarged cell somas and dendritic arbors (Morillo et al. 2010), suggesting that somatic tetraploidy is associated with morphological and functional diversity in the nervous system.

# 4 p75<sup>NTR</sup> in Neurodegeneration: The Tetraploid Connection

Cell cycle reentry in neurons represents a critical feature of both acute neuronal damage and chronic neurodegenerative diseases (Wang et al. 2009). In this regard, cell cycle regulators have been proposed as therapeutic targets for stroke (Osuga et al. 2000), excitotoxicity (Verdaguer et al. 2004), trauma (Di Giovanni et al. 2005), and Alzheimer's disease (Woods et al. 2007).

In cases of acute injury to the CNS such as kainic acid excitotoxicity and trauma, cell cycle reentry is usually stopped at the G1/S phase and it leads to rapid apoptosis (Kuan et al. 2004; Byrnes et al. 2007). In contrast, chronic neurodegenerative diseases are normally associated with DNA synthesis in particular neuronal populations, which can be arrested at the G2/M transition and survive for some time before dying by apoptosis. This is the case, for instance, of neurons subjected to ischemia-hypoxia (Byrnes et al. 2007; Burns et al. 2007).

These neurons incorporate BrdU and remain alive for days, before they die. Examples of cell cycle reentry and hyperploidization have also been shown to occur in Alzheimer's disease (Yang et al. 2001; Mosch et al. 2007; Arendt et al. 2010). Furthermore, increased expression in neurons of cell cycle regulators involved in G1/S transition has been described in patients suffering amyotrophic lateral sclerosis (Ranganathan and Bowser 2003, 2010; Ferraiuolo et al. 2007), Parkinson's disease (Höglinger et al. 2007), Huntington's disease (Pelegrí et al. 2008; Fernandez-Fernandez et al. 2011), and different tauopathies (Stone et al. 2011). Furthermore, the Parkin2 gene, known to be a frequent cause of early-onset Parkinson's disease, is a tumor suppressor gene whose inactivation results in an increase in cyclin E levels (Veeriah et al. 2010). Neuronal cell cycle reentry in pathological states has often been interpreted as de novo adult neurogenesis (Fallon et al. 2000; Arvidsson et al. 2002; Curtis et al. 2003; Becker et al. 2007). Nevertheless, this conclusion has been proved to be erroneous in some cases. For instance, the use of conditional transgenic mice expressing EGFP under the control of the nestin promoter allowed Burns et al. (2007) to demonstrate that ischemiahypoxia-dependent BrdU incorporation mostly occurs in differentiated neurons.

In the adult nervous system, p75<sup>NTR</sup> is reexpressed in various neuropathological conditions, including Alzheimer's disease (Hu et al. 2002), amyotrophic lateral sclerosis (Lowry et al. 2001), Parkinson's disease (Chen et al. 2008), and Huntington's disease (Zuccato et al. 2008), likely participating in the etiology of neurodegeneration (Dechant and Barde 2002). p75<sup>NTR</sup> is therefore a candidate for the induction of cell cycle reentry leading to tetraploidy in the affected neurons (Frade and López-Sánchez 2010). Since developmentally regulated tetraploidy in neurons correlates with cell body and dendritic tree enlargement (see above), it is conceivable that a similar effect may occur in adult neurons undergoing de novo tetraploidization, thus explaining, at least partially, the pathological signs of the disease. Structural alterations in neurons becoming tetraploid may correlate with functional changes, including alterations of metabolic activity, higher rate of membrane biosynthesis, changes in neuronal circuits, and alterations in electrical signal propagation (see Frade and López-Sánchez 2010). Furthermore, genome duplication may result in changes of gene expression in particular loci, as those observed in liver and heart cells (Anatskaya and Vinogradov 2007) and tetraploid mouse embryos (Kawaguchi et al. 2009). It is likely that all these alterations associated with tetraploidy in neurons could functionally compromise the affected neurons. Further insights on the molecular mechanisms triggering tetraploidy in neurons will probably be crucial for the design of appropriate therapeutic tools for patients that suffer from neurodegenerative diseases.

### 5 Conclusion

 $p75^{NTR}$  is a pleiotropic receptor that fulfills several critical functions in the nervous system, including the regulation of the cell cycle in both embryonic and adult nervous system. Indeed,  $p75^{NTR}$  seems to be involved in stem cell

maintenance, neurogenesis, and neuronal tetraploidization. The molecular mechanisms used by p75<sup>NTR</sup> to regulate the cell cycle are far from being fully understood, but they are likely based on its capacity to interact with a number of cell cycle regulators, as well as its ability to translocate its intracellular domain to the nucleus to act as a transcriptional regulator. The capacity of p75<sup>NTR</sup> to regulate the cell cycle has opened new research avenues not only focused on the normal development of the vertebrate nervous system but also related with neuropathological conditions, since deregulation of this receptor in the adult brain could participate in neurodegenerative diseases. It is likely that future research in this field will yield hints of possible therapeutic approaches for these devastating diseases.

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# p75NTR Processing and Signaling: Functional Role

# Ramiro D. Almeida and Carlos B. Duarte

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### Abstract

The p75 neurotrophin receptor  $(p75^{NTR})$  was the first neurotrophin receptor to be identified and, more than two decades after, significant advances have been made in understanding its biological actions.  $p75^{NTR}$  belongs to the tumor necrosis factor receptor superfamily (TNFRS) and is particularly important during developmental neural cell death and under injury conditions in the adulthood. However, recent studies shed new light into the functions of  $p75^{NTR}$  in the nervous system.  $p75^{NTR}$  either acting autonomously or as part of a complex with other co-receptors regulates neuronal survival and differentiation, cell cycle and proliferation, axonal outgrowth, and synaptic

R.D. Almeida

Center for Neuroscience and Cell Biology (CNC), University of Coimbra, Coimbra, Portugal

C.B. Duarte (⊠)

Center for Neuroscience and Cell Biology (CNC) and Department of Life Sciences, University of Coimbra, Coimbra, Portugal e-mail: cbduarte@ci.uc.pt

plasticity. Here, we will review the main findings of the last two decades as well as new findings that reveal additional and unexpected roles for p75<sup>NTR</sup>.

Keywords

Axonal growth • Cell cycle • Neuronal death • Neurotrophins • p75NTR • Synaptic plasticity

### 1 Introduction

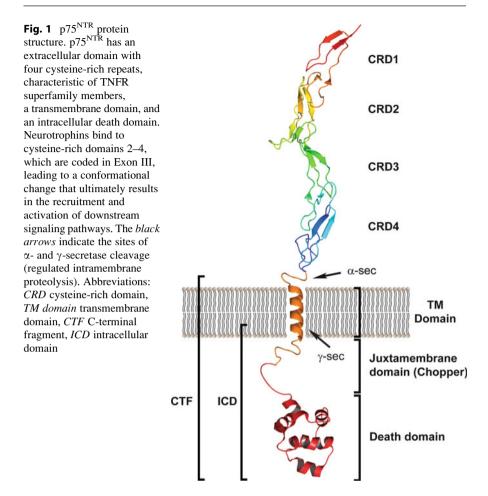
The p75 neurotrophin receptor  $(p75^{NTR})$  belongs to the tumor necrosis factor receptor superfamily (TNFRS) and was first identified by Shooter and colleagues (reviewed in Herrup and Shooter 1973).  $p75^{NTR}$  has an important role in the development of the nervous system and accordingly is mainly expressed during development, decreasing in the adulthood, except in a few selected populations of neurons and under pathological conditions, when it is re-expressed (reviewed in Dechant and Barde 2002). It should also be emphasized that  $p75^{NTR}$  has critical roles in glial biology and outside the nervous system, namely, in the immune and vascular systems.

All known mammalian neurotrophins, nerve growth factor (NGF), brainderived neurotrophic factor (BDNF), neurotrophin-3 (NT-3), and neurotrophin 4/5 (NT-4/5), bind to p75<sup>NTR</sup> with similar affinities. However, they bind their cognate tropomyosin-related kinase (Trk) receptor with different specificities. Generally, NGF binds TrkA, BDNF and NT-4/5 activate TrkB, and NT-3 recognizes TrkC, and this selectivity is believed to promote neuronal survival and differentiation in distinct populations of neurons (reviewed in Lu et al. 2005). Neurotrophins can also induce cell death by activating p75<sup>NTR</sup> and this causes an apparent contradiction. The latter effects are mediated by proneurotrophins, the immature uncleaved form, which bind  $p75^{NTR}$  with relatively higher affinity than to Trk receptors, thereby promoting cell death. Accordingly, the mature forms of neurotrophins preferentially bind Trk receptors (Lee et al. 2001; Teng et al. 2005). This specificity is achieved through the formation of molecular complexes with different co-receptors. While mature neurotrophins bind to a complex of p75<sup>NTR</sup> and Trk, the proforms preferentially bind to a complex of p75<sup>NTR</sup> and Sortilin, another type I transmembrane protein, but structurally unrelated to p75<sup>NTR</sup> (reviewed in Nykjaer et al. 2005). This feature reveals a distinctive characteristic of p75<sup>NTR</sup>: signaling independently and in conjunction with its co-receptors. The best studied p75<sup>NTR</sup> partner is the Trk receptor, and the molecular basis of their interaction has been extensively studied. p75<sup>NTR</sup> is able to form complexes with TrkA, TrkB, and TrkC, altering its affinity and signaling properties (reviewed in Chao 2003). Recent studies revealed new roles for p75<sup>NTR</sup> in the nervous system and along with it new co-receptors and cytosolic interactors were described. In fact, p75<sup>NTR</sup> was described to regulate additional aspects of neuronal development like axonal growth, cell cycle, and synaptic plasticity (reviewed in Ibanez and Simi 2012). Although exciting, these novel observations raise the question of how  $p75^{NTR}$  signaling specificity is achieved with a relatively high number of signaling partners. It is believed that such specificity of  $p75^{NTR}$  actions is likely to be dependent on the developmental and cellular context in which the receptor is integrated.

### 2 Expression: Isoforms and Co-receptors

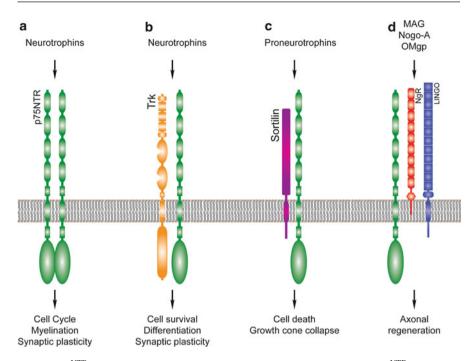
The p75<sup>NTR</sup> is widely expressed in the nervous system during development. In the developing CNS, the  $p75^{NTR}$  is present in spinal motor neurons and brain stem motor nuclei, lateral geniculate nucleus, medial terminal nucleus of the accessory optic tract, ventral and dorsal cochlear nucleus, thalamic nucleus, nucleus of the lateral lemniscus, amygdala, cortical subplate neurons, olivary pretectal nucleus, cuneate nucleus, and gracile nucleus, in addition to Purkinje neurons and the external granule layer and deep nuclei of the cerebellum (Buck et al. 1988; Ernfors et al. 1988; Heuer et al. 1990; Schatteman et al. 1988). In the adult nervous system, the p75<sup>NTR</sup> is found mainly within the basal forebrain (Hefti et al. 1986; Schatteman et al. 1988; Springer et al. 1987; Yan and Johnson 1988), and lower expression levels are observed in the caudate/putamen neurons (Henry et al. 1994), cerebellar Purkinje neurons (Cohen-Cory et al. 1991; Koh et al. 1989; Mufson et al. 1991; Shelton and Reichardt 1986), motor neurons (Armstrong et al. 1991; Ernfors et al. 1989), and within several central nuclei (Cohen-Cory et al. 1991; Koh et al. 1989; Schatteman et al. 1988; Shelton and Reichardt 1986; Sofroniew et al. 1989). p75<sup>NTR</sup> is expressed in neurons and glia (Cragnolini and Friedman 2008), and their expression is induced by many types of injury, including transient forebrain or global cerebral ischemia (Junier et al. 1994; Lee et al. 1995; Oderfeld-Nowak et al. 2003), in ischemia induced by cortical devascularization (Angelo et al. 2009), and in the brain of rats subjected to pilocarpine-induced seizures (Roux et al. 1999; Unsain et al. 2008) [for a detailed review see Ibanez and Simi (2012)]. In the peripheral nervous system, the p75<sup>NTR</sup> is expressed in dorsal root ganglia sensory neurons as well as in a subpopulation of enteric and parasympathetic neurons (Carroll et al. 1992; Schatteman et al. 1993; Verge et al. 1992; Yan and Johnson 1988). Axotomy of peripheral motor neurons, resembling the lesions resulting from severe traumatic injuries of the limbs, was also shown to upregulate p75<sup>NTR</sup> expression in adult animals (Ferri et al. 2002). Furthermore, p75<sup>NTR</sup> is present outside the nervous system, where the highest expression of the receptor is found (reviewed in Barker 1998).

The p75<sup>NTR</sup> belongs to the tumor necrosis factor family of receptors (TNFR), comprising an extracellular region with four cysteine-rich domains, each with six cysteine residues, a single transmembrane domain, and the intracellular region. The latter region contains two domains that are able to mediate intracellular signaling, the Chopper domain and the death domain homologous to that present in the TNFR family (reviewed in Skeldal et al. 2011). The Chopper domain is a juxtamembrane region with the ability to induce cell death when bound to the membrane (Coulson et al. 2000; Underwood et al. 2008) (Fig. 1). The p75<sup>NTR</sup> can bind all four neurotrophins (NGF, BDNF, NT3, and NT4/NT5) (Rodriguez-Tebar et al. 1992;



Ryden et al. 1995; Urfer et al. 1994), as well as to proNGF and proBDNF (Lee et al. 2001; Teng et al. 2005). The crystal structure of the extracellular domain of  $p75^{NTR}$  complexed with NGF (Aurikko et al. 2005) or NT3 (Gong et al. 2008) showed a symmetrical 2:2 receptor-ligand complex. However, a different study reported that  $p75^{NTR}$  binds NGF in a 1:2 stoichiometry (He and Garcia 2004), which may correspond to an intermediate step before a symmetrical 2:2 receptor-ligand complex is formed. The  $p75^{NTR}$  forms dimers even in the absence of neurotrophins through cysteinyl residues within the transmembrane domain, and it was proposed that the interaction of the neurotrophins with the receptor complex induces a conformational change in the  $p75^{NTR}$  intracellular domain which may allow the recruitment of the effector molecules NRIF and TRAF6 (Vilar et al. 2009a, b). The activation of  $p75^{NTR}$  by all neurotrophins contrasts with the more stringent specificity of the Trk family of receptors for neurotrophins.

The biological effects of proNGF and proBDNF mediated by activation of  $p75^{NTR}$  require the co-receptor sortilin which interacts with the prodomain of the



**Fig. 2** p75<sup>NTR</sup> co-receptor complexes and their biological function. To date, p75<sup>NTR</sup> was shown to interact with a diversity of receptors, namely, Sortilin, LINGO-1, NgR, and Trk receptors. The outcome of these interactions is a variety of cellular responses including cell death, synaptic plasticity, neurite outgrowth and cell cycle regulation. The C-terminal fragment of p75<sup>NTR</sup> can also bind other cytosolic interactors (not depicted here), increasing the diversity and complexity of signaling mechanisms that are regulated by this receptor. Abbreviations: *MAG* myelin-associated glycoprotein, *OMGP* oligodendrocyte myelin glycoprotein

neurotrophins (Nykjaer et al. 2004; Teng et al. 2005) (Fig. 2c). However, it is still not yet clear whether each prodomain of the dimeric neurotrophin molecules binds to a different sortilin molecule or whether two prodomains bind to the same sortilin protein (Feng et al. 2010). During development, sortilin is expressed in the brain cortex, hippocampus, and neural retina, and this pattern is preserved in the adulthood, when the protein is particularly abundant in pyramidal cells of the hippocampus (Hermans-Borgmeyer et al. 1999; Sarret et al. 2003). Activation of the p75<sup>NTR</sup>/sortilin complex by proNGF or proBFNF was shown to induce apoptotic cell death in cultured sympathetic neurons, but this effect was not observed in cells derived from mice lacking p75<sup>NTR</sup> or in the presence of the sortilin antagonist neurotensin (Nykjaer et al. 2004; Teng et al. 2005).

A truncated isoform of p75<sup>NTR</sup> has been described, resulting from alternative splicing and lacking the cysteine-rich repeats 2, 3 and 4 (s-p75<sup>NTR</sup>) (Paul et al. 2004; von Schack et al. 2001), which are required for neurotrophin binding (Aurikko et al. 2005; Feng et al. 2010; Gong et al. 2008; He and Garcia 2004). Although these variants are unable to bind neurotrophins, they may regulate cellular activity by

interacting with some of the intracellular signaling proteins regulated by the fulllength p75<sup>NTR</sup>. Furthermore, a neurotrophin receptor homologue-2 (NRH2, also termed PLAIDD or NRADD) was described, which lacks the ligand-binding domain present in the p75<sup>NTR</sup> but has some similarity with its transmembrane, juxtamembrane and the death domain regions (Kanning et al. 2003; Murray et al. 2004). The expression of NRH2 receptors is developmentally regulated, decreasing in adult tissues (Frankowski et al. 2002; Wang et al. 2003). Although the homology between p75<sup>NTR</sup> and NRH2 suggests that it may act as a functional analogue of the s-p75<sup>NTR</sup>, the differences between the intracellular domains of the two receptor proteins may provide some specificity in the biological activity of NRH2 receptors (Kim and Hempstead 2009). NRH2 facilitates the formation of a complex between p75<sup>NTR</sup> and sortilin, and was proposed to act by impairing the degradation of sortilin in the lysosomes and by redistributing sortilin to the surface (Kim and Hempstead 2009). However, additional studies are required to determine the stoichiometry of the interactions between p75<sup>NTR</sup>, sortilin, and NRH2.

In addition to its signaling activity, the  $p75^{NTR}$  may also interact with the TrkA (Huber and Chao 1995), TrkB, and TrkC receptors (Bibel et al. 1999), thereby modulating their activity. In particular, the  $p75^{NTR}$ /TrkA complexes show an increased affinity for NGF when compared with each one of the receptors expressed independently (Hempstead et al. 1991), and the co-expression of the two classes of receptors allows TrkA-mediated responses at lower concentrations of NGF (Horton et al. 1997; Lee et al. 1994). The TrkA and  $p75^{NTR}$  were shown to interact through their intracellular domains even before NGF stimulation, and this association may contribute to the observed high-affinity NGF-binding sites (Iacaruso et al. 2011).

# 3 p75<sup>NTR</sup> Proteolysis and Signaling in Cell Death and Survival

The absence of catalytic activity in the p75<sup>NTR</sup> intracellular region suggests that the intracellular signaling activity is mediated by binding partners that may be recruited upon receptor activation and/or constitutively bound. In fact, more than 20 different intracellular binding partners of p75<sup>NTR</sup> were identified up to now (Barker 2004; Bronfman and Fainzilber 2004; Dechant and Barde 2002; Roux and Barker 2002; Schecterson and Bothwell 2010).

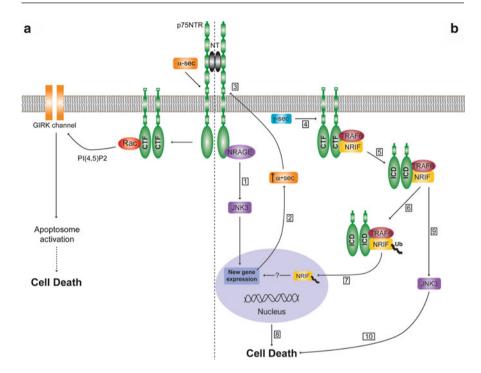
Initial studies showed a role for p75<sup>NTR</sup> in the facilitation of survival signals induced by Trk receptors, activated by the mature form of neurotrophins, and this effect has been attributed to an increase in the affinity of the Trk receptor for the ligand (Roux and Barker 2002) (Fig. 2b). Furthermore, p75<sup>NTR</sup> increases the specificity of Trk receptors for particular ligands, which may change the pattern of signaling response to stimulation by each neurotrophin (Clary and Reichardt 1994). The mechanism whereby p75<sup>NTR</sup> increases Trk receptor activity has not been elucidated, but should involve the intracellular region of the receptor since extracellular domain mutations of p75<sup>NTR</sup> proteins also allow the formation of

high-affinity Trk binding sites (Esposito et al. 2001). This is supported by results obtained with Förster resonance energy transfer showing that p75<sup>NTR</sup> and TrkA receptors physically interact through their intracellular domains, even before stimulation with NGF (Iacaruso et al. 2011). A model was proposed according to which Trk receptor activation leads to activation of the transmembrane cysteine protease  $\alpha$ -secretase tumor necrosis factor- $\alpha$  converting enzyme/a disintegrin and metallopeptidase domain (TACE/ADAM) 17 by MEKdependent phosphorylation, which is followed by p75<sup>NTR</sup> cleavage. TACE/ ADAM17 displays  $\alpha$ -secretase activity, cleaving p75<sup>NTR</sup> in the extracellular juxtamembrane domain, releasing the extracellular domain and giving rise to a product that remains anchored at the membrane. According to this model, the  $p75^{NTR}$  cleavage product feeds back to enhance Trk-induced neurotrophin signaling, thereby increasing ERK and Akt activity which enhance cell survival (Ceni et al. 2010; Kommaddi et al. 2011). Whether the p75<sup>NTR</sup> cleavage products directly interact with Trk receptors to increase the affinity for ligands remains to be elucidated.

The  $p75^{NTR}$  may also be coupled to cell death when activated by proneurotrophins, and in this case, sortilin has been shown to act as a co-receptor, increasing the affinity of  $p75^{NTR}$  for the ligand (Nykjaer et al. 2004). Sortilin interacts with the prodomain of neurotrophins, and the signaling that leads to cell death may occur in the absence or in the presence of Trk receptors. The role of  $p75^{NTR}$  and sortilin in proneurotrophin-induced cell death is supported by studies showing that antibodies against sortilin,  $p75^{NTR}$ , or proNGF protect cells from apoptotic death (Beattie et al. 2002; Nykjaer et al. 2004; Volosin et al. 2006, 2008). Furthermore, an excess of the prodomain of proNGF or neurotensin, which also binds sortilin (Quistgaard et al. 2009), inhibits proNGF-induced cell death (Jansen et al. 2007; Nykjaer et al. 2004; Rogers et al. 2010; Volosin et al. 2006).

An interesting feature of p75<sup>NTR</sup> function is its ability to undergo proteolysis with the generation of fragments with signaling capability. p75<sup>NTR</sup> proteolysis was described to be required for cell survival (addressed below), cell migration, neurite growth inhibition, and cell cycle. In fact, several intracellular domain interactors that regulate cell cycle, like Schwann cell factor-1 (Sc-1), brain-expressed X-linked 1 (Bex1), and melanoma antigen gene (MAGE), have been found in the nucleus after p75<sup>NTR</sup> activation (reviewed in Skeldal et al. 2011), but p75<sup>NTR</sup> regulated intramembrane proteolysis (RIP) has been best studied in neuronal death. The process of p75<sup>NTR</sup> proteolytic cleavage occurs through the sequential action of two distinct proteins. First, the extracellular domain (ECD) of p75<sup>NTR</sup> is cleaved by TACE/ADAM17, generating a membrane-bound C-terminal fragment (CTF). The CTF undergoes a second cleavage by the presentlin-dependent  $\gamma$ -secretase, releasing the intracellular domain (ICD) into the cytoplasm (Jung et al. 2003; Kanning et al. 2003) (Fig. 3). NRH2 can also undergo proteolytic cleavage, releasing its cytoplasmic domain, which unlike the ICD has higher stability (Kanning et al. 2003).

The  $p75^{NTR}$  ICD regulates neuronal cell death in vitro and in vivo. Neurotrophin binding to  $p75^{NTR}$  was shown to promote  $\gamma$ -secretase-dependent



**Fig. 3**  $p75^{NTR}$  proteolysis and cell death signaling pathways. Regulated intracellular proteolysis of  $p75^{NTR}$  leading to neuronal cell death can act through two distinct pathways. (a) In one model, the 29 amino acid intracellular juxtamembrane region of  $p75^{NTR}$  named "chopper" induces cell death through activation of G-protein-coupled inwardly rectifying potassium (GIRK/Kir3) channels. The corresponding potassium efflux supposedly leads to apoptosome activation and by consequence neuronal cell death. (b) In another model  $p75^{NTR}$  activates JNK 3 (1), which leads to the transcription-dependent upregulation of  $\alpha$ -secretase (2) that cleaves  $p75^{NTR}$  extracellular domain (3).  $\gamma$ -secretase releases the ICD into the cytoplasm (4) which associates with the DNA-binding protein NRIF and the E3 ubiquitin ligase TRAF6 (5). The formation of this ternary complex leads to TRAF6-dependent ubiquitination of NRIF (6) and nuclear translocation (7), a process required for  $p75^{NTR}$ -dependent cell death (8). In parallel, this ternary complex further activates JNK3 (9), amplifying the apoptotic signal (10). Abbreviations: *JNK3* c-Jun N-terminal kinase 3, *NRAGE* neurotrophin-receptor-interacting MAGE (melanoma-associated antigen) homologue

cleavage, with the cytosolic release of ICD and subsequent nuclear translocation (Frade 2005; Kenchappa et al. 2006). The complete picture is however considerably more complicated and involves a dual-phase signaling mechanism. Stimulation of the receptor with BDNF induces the activation of JNK3, leading to the upregulation of TACE/ADAM17 by a transcription-dependent mechanism (Kenchappa et al. 2010). The p75<sup>NTR</sup> is first cleaved by TACE/ADAM17 and then by the  $\gamma$ -secretase complex, releasing the intracellular domain (ICD) of the receptor and allowing the translocation of neurotrophin-receptor-interacting factor (NRIF) to the nucleus. The p75<sup>NTR</sup> ICD can also translocate to the nucleus

where it regulates gene expression (Parkhurst et al. 2010). The nuclear translocation of NRIF depends on its polyubiquitination mediated by the E3 ubiquitin ligase TRAF6, and is necessary for p75<sup>NTR</sup>-mediated apoptosis (Geetha et al. 2005; Kenchappa et al. 2006). In addition, the ICD/TRAF6/NRIF complex can activate JNK3 directly further amplifying the cell death cascade (Fig. 3b). NRIF nuclear translocation seems to be of particular importance in the period of naturally occurring superior cervical ganglion (SCG) neuron death since this protein is observed in the nucleus of SCG neurons at postnatal day 4 but not at postnatal day 24, a developmental time point when neuron elimination has diminished (Kenchappa et al. 2010). How the nuclear translocation of NRIF induces apoptosis has not yet been elucidated, but considering that NRIF is a zinc finger DNA-binding protein, it may act through altering the expression of proapoptotic genes (Kenchappa et al. 2006). NRIF was also shown to mediate the apoptotic signaling initiated by p75<sup>NTR</sup> in hippocampal neurons after pilocarpineinduced seizures (Volosin et al. 2008). The protein NRAGE (neurotrophinreceptor-interacting MAGE homolog), a member of the MAGE (melanoma antigen) family, also interacts with the p75<sup>NTR</sup> intracellular region, facilitating the receptorinduced apoptotic death (Salehi et al. 2000). The effects of NRAGE are mediated by stimulation of JNK and c-Jun, which is followed by the cytochrome C release from the mitochondria, and activation of caspases-9, -3, and -7 to induce cell death (Salehi et al. 2002).

Under certain conditions, the C-terminal fragment of p75<sup>NTR</sup> resulting from metalloprotease cleavage may induce cell death without additional cleavage by  $\gamma$ -secretase to generate the ICD (Coulson et al. 2000, 2008; Underwood et al. 2008). The p75<sup>NTR</sup> CTF contains a cytoplasmic juxtamembrane sequence of 29 amino acids that is both necessary and sufficient to trigger cell death (Coulson et al. 2000). According to the model proposed, the cleavage of p75<sup>NTR</sup> followed by palmitoylation of the CTF causes its translocation to cholesterol-rich domains of the plasma membrane (Underwood et al. 2008). The p75<sup>NTR</sup> CTF upregulates the production of phosphatidylinositol 4,5-bisphosphate (PtdIns(4,5)P<sub>2</sub>) through activation of Rac, and the locally generated PtdIns(4,5)P<sub>2</sub> activate G-protein-coupled inwardly rectifying potassium (GIRK/Kir3) channels (Coulson et al. 2008) (Fig. 3a). The resulting increase in  $K^+$  permeability with consequent reduction in the  $[K^+]_i$  may contribute to induce the apoptotic pathway. However, additional studies are required to determine under which conditions this pathway is active in vivo and how the CTF- and the ICD-mediated pathways are differentially regulated.

In addition to the activation of apoptotic signaling pathways, p75<sup>NTR</sup> may also be coupled to the induction of cellular alterations that prevent the trophic effects of Trk neurotrophin receptors, thereby contributing to cell death. Such a mechanism was described in basal forebrain neurons, one of the few CNS populations that express p75<sup>NTR</sup> throughout life. Stimulation of p75<sup>NTR</sup> with proNGF upregulates PTEN (phosphatase and tensin homolog deleted on chromosome 10) in basal forebrain neurons, a protein known to antagonize the activation of Akt by PI3-K. When the cells are stimulated simultaneously with proNGF and BDNF, to activate  $p75^{NTR}$  and TrkB receptors, respectively, the inhibition of the TrkB-induced PI3-K signaling by PTEN is sufficient to abrogate the trophic effects of TrkB signaling (Song et al. 2010). Accordingly, inhibition of PTEN was shown to prevent neuronal loss in the hippocampus following pilocarpine-induced seizures, a model of  $p75^{NTR}$ -mediated neuronal death (Song et al. 2010).

## 4 Axonal Growth

The antagonistic functions of neurotrophins and their receptors are well depicted in sympathetic neuron growth and target innervations. NGF acting through TrkA induces axonal growth, while BDNF acting on  $p75^{NTR}$  inhibits it (Kohn et al. 1999). These evidences are supported in vivo where the pineal gland, a sympathetic target organ, of BDNF –/– mice is hyperinnervated (Kohn et al. 1999). This axonal pruning depends on activity-dependent release of BDNF by the wining axons that in turn acts on  $p75^{NTR}$  present in the losing axons, antagonizing NGF/TrkA (Singh and Miller 2005; Singh et al. 2008). This model provides an elegant explanation of how competing axons can promote growth inhibition and axonal degeneration on less favored neighboring axons.

Currently, it is widely accepted that central nervous system (CNS) neurons do not have the ability to regenerate upon injury. Several factors account for this phenomenon such as the scar generation at the injury site, reduced regrowth capability of mature neurons, and myelin-associated inhibitory factors (MAIFs) (Chen et al. 2009). Myelin-derived molecules such as myelin-associated glycoprotein (MAG), oligodendrocyte myelin glycoprotein (oMgp), and Nogo exert their action through a common receptor, the Nogo receptor (NgR) (Fournier et al. 2001), and this effect is mediated by activation of a small G protein, the Ras homologue member A (RhoA) (Niederost et al. 2002; Yamashita et al. 2002). However, NgR is a glycosyl-phosphatidylinositol (GPI)-linked membrane protein, with no cytoplasmic domain and therefore, it is not capable of transducing an appropriate signal upon ligand binding. Subsequently, it was demonstrated that p75<sup>NTR</sup> is a coreceptor for NgR (Wang et al. 2002; Wong et al. 2002). Neurons from p75<sup>NTR</sup> knockout mice are no longer responsive to myelin, and blocking NgR-p75<sup>NTR</sup> interaction has a similar effect (Wang et al. 2002). However, in a reconstituted nonneuronal system, NgR and p75<sup>NTR</sup> alone are not able to activate RhoA, indicating that an additional molecule or molecules are required. Exogenous expression of a dominant negative LINGO-1 in cerebellar granular neurons reduced myelin inhibition (Mi et al. 2004) (Fig. 2d), indicating LINGO-1 forms a signaling complex with NgR and p75<sup>NTR</sup>.

p75<sup>NTR</sup> directly regulates RhoA activation. Barde and colleagues demonstrated that neurotrophin binding to p75<sup>NTR</sup> inhibits RhoA activation, increasing the rate of neurite outgrowth (Yamashita et al. 1999). A similar effect is observed on filopodial dynamics where neurotrophin treatment of retinal or dorsal root ganglion (DRG) neurons leads to a decrease in RhoA activity, which in turns increases filopodial length. On the other hand, when MAIFs bind to p75<sup>NTR</sup>, RhoA is activated and

neurite outgrowth reduced. This effect is mediated by the Rho-GDP dissociation inhibitor (Rho-GDI) which interacts with the cytoplasmic domain of  $p75^{NTR}$ . Usually, RhoA is bound to Rho-GDI, but the MAG or Nogo proteins facilitate the interaction between  $p75^{NTR}$  and Rho-GDI, releasing RhoA (Yamashita and Tohyama 2003), which then leads to actin depolymerization. Additionally, MAG induces  $\alpha$ - and then  $\gamma$ -secretase-mediated proteolysis of  $p75^{NTR}$  and this cleavage is required for the activation of RhoA and neurite outgrowth inhibition (Domeniconi et al. 2005). However,  $p75^{NTR}$  is only expressed in selected populations of mature neurons, raising the question of how myelin inhibitory signals are transduced in  $p75^{NTR}$ -lacking cells. TROY/TAJ, another TNFR homologue, broadly expressed in postnatal and adult neurons, can also bind NgR and LINGO-1, to form a functional complex and activate RhoA in the presence of myelin inhibitors (Park et al. 2005; Shao et al. 2005). Together, these studies established that  $p75^{NTR}$  has a crucial role in axonal regeneration.

It is puzzling why the nervous system would increase the expression of  $p75^{NTR}$  when its effects are so deleterious to neurons. Currently, there is no clear answer to this question, but we can speculate that the detrimental effects of  $p75^{NTR}$  in neuronal injury could be due to a deregulation of a homeostatic mechanism (reviewed in Ibanez and Simi 2012). In line with this, a recent study in the uninjured adult mice brain showed that myelin acting through  $p75^{NTR}$  is responsible for the axonal degeneration of aberrant septal cholinergic axons projecting to the cerebral cortex and hippocampus through the supra-callosal pathway (Park et al. 2010). Interestingly, this effect required BDNF binding to  $p75^{NTR}$  since an antibody against BDNF rescued myelin-induced inhibition of axonal growth and axonal degeneration (Park et al. 2010). These results provide an explanation to the intrinsic inability of CNS neurons to regrow on myelin inhibitory substrates.

p75<sup>NTR</sup> also regulates growth cone collapse. Recently, it was shown that proNGF induces growth cone collapse in cultured hippocampal neurons. Binding of proNGF to a p75<sup>NTR</sup> and sortilin-related VPS10 domain-containing receptor 2 (SorCS2) complex decreases Rac activity and inactivates fascin, an actin-bundling protein, regulating actin dynamics (Deinhardt et al. 2011). A single nucleotide polymorphism (SNP) is present on the 5' proBDNF sequence at codon 66, producing an amino acid substitution (Val66Met). In humans, the Met66BDNF was associated with abnormal hippocampal activation and poorer episodic memory (Egan et al. 2003). Interestingly, preliminary studies suggest that recombinant Met66 but not Val66 BDNF prodomain induces growth cone collapse in cultured hippocampal neurons and this effect is dependent on p75<sup>NTR</sup> and SorCS2 (Anastasia et al. 2013). The authors claim Met66 prodomain is intrinsically disordered with some transient secondary structure. However, it is not clear how proNGF and proBDNF(Val66), which share a high degree of homology in their prodomain, elicit different results.

Recently, p75<sup>NTR</sup> has been involved in two other systems of axonal guidance, Sema3A and ephrin-A. Eph/ephrins are membrane proteins that regulate neuronal migration, synapse formation, and axon guidance. The eph/ephrin system is divided into class A and B, and they can signal bidirectionally; eph signaling is considered forward while ephrin signaling is considered reverse (reviewed in Xu and Henkemeyer 2012). Similarly to the Nogo receptor, class A ephrins are GPI-linked proteins and require a co-receptor as a transducing unit. Using retinal ganglion cells (RGCs), a widely used axon guidance model system, O'Leary and colleagues reported that ephrin-As colocalize with p75<sup>NTR</sup> in retinal axons and this association is required for Fyn phosphorylation upon ephA binding (Lim et al. 2008). Moreover, this requirement is specific to ephrin reverse signaling since eph forward signaling is not affected (Lim et al. 2008). Sema3A is a secreted protein that regulates growth cone collapse and repulsion, and signals through plexins and neuropilins (Pasterkamp 2012). Sema3A was found to increase the co-localization of p75<sup>NTR</sup> and neuropilin-1 in DRG growth cones, and inhibits the direct association of plexinA4/neuropilin-1 complex leads to increased sensitivity of DRG neurons from p75<sup>NTR</sup> (ExonIII-/-) mice to Sema3A (Ben-Zvi et al. 2007).

## 5 Regulation of Cell Cycle

The p75<sup>NTR</sup> is expressed very early during development of the nervous system (Yan and Johnson 1987, 1988), when many precursor cells are still not differentiated and before targeting specific neuronal populations for programmed cell death. The role of p75<sup>NTR</sup> signaling in the modulation of cell cycle progression has been investigated in vitro and in vivo. Thus, the pheochromocytoma PC12 cells differentiate in response to NGF when stimulated in the G1 phase (but not during the other phases of the cell cycle) (Rudkin et al. 1989), and this effect correlates with the transient surface expression of p75<sup>NTR</sup> (Ito et al. 2002, 2003; Urdiales et al. 1998). Furthermore, stimulation of p75<sup>NTR</sup> with NGF was shown to prevent the progression through the G1 phase of the cell cycle in astrocytes (Cragnolini et al. 2009; Cragnolini et al. 2012). Activation of p75<sup>NTR</sup> was also shown to play a role in BDNF-induced differentiation of neural precursor cells in culture (Hosomi et al. 2003) and may be involved in the development of dendrites by purified subventricular cells cultured in the presence of neurotrophins (Gascon et al. 2005).

In contrast with the differentiating effects described above, p75<sup>NTR</sup> receptors were shown to mediate the NGF-induced proliferation of mouse embryonic stem cells in culture (Moscatelli et al. 2009). p75<sup>NTR</sup> is also expressed in adult dentate gyrus progenitor cells, playing a role in the proliferation and/or early maturation of neuronal and glial cells, as well as other cell types (Bernabeu and Longo 2010). Furthermore, in vitro and in vivo studies identified a population of highly proliferative subventricular zone cells in the adult rat, characterized by the expression of p75<sup>NTR</sup> and showing an increased rate of neuroblast generation in response to BDNF (Giuliani et al. 2004; Young et al. 2007). However, under disease or injury conditions, the potential of these p75<sup>NTR</sup>-expressing cells present in the subventricular zone may be altered, to yield glial progenitors in addition to their neurogenic potential. In patients and in animal models of multiple sclerosis,

the p75<sup>NTR</sup>-positive cells in the subventricular zone may give rise to oligodendrocyte progenitors (Petratos et al. 2004). The amyloid-beta peptide (AB), which is accumulated in amyloid plaques and generally recognized as a key feature of Alzheimer's disease, was also shown to stimulate adult SVZ neurogenesis, by a mechanism dependent on the expression of p75<sup>NTR</sup> by the precursor cells and activation of the receptors through cleavage by metalloproteases (Sotthibundhu et al. 2009). This correlates with the results showing an interaction of AB with  $p75^{NTR}$ (Yaar et al. 1997, 2002), but was observed mainly in younger animals. Based on these observations, it was suggested that overstimulation of p75<sup>NTR</sup>-positive progenitor cells early in life may lead to the depletion of a pool of stem cells with a consequent downregulation in basal neurogenesis (Sotthibundhu et al. 2009). Excitotoxic stimulation of the adult striatum also induces the expression of the p75<sup>NTR</sup> protein in astrocytic progenitor cells (Hanbury et al. 2002), and pilocarpineinduced seizures upregulated  $p75^{NTR}$  protein levels in neurons and astrocytes (Cragnolini et al. 2009; Cragnolini et al. 2012). Since p75<sup>NTR</sup> controls the astrocytic proliferation (see above) and its ligand (NGF) is produced by astrocytes under inflammatory conditions (Friedman et al. 1996; Oderfeld-Nowak and Bacia 1994), NGF may have an autocrine or paracrine effect in the regulation of reactive gliosis in vivo after injury.

The diversity of mechanisms contributing to the differential effects of p75<sup>NTR</sup> and NGF on cell cycle is not fully understood, but these responses are likely to be mediated by the interaction of the receptor with different intracellular proteins under distinct cellular contexts. Several of the p75<sup>NTR</sup> binding partners are regulators of the cell cycle, and their interaction with the receptors is controlled by ligand binding. The zinc finger protein SC1 interacts with the intracellular region of the p75<sup>NTR</sup>, being translocated to the nucleus following stimulation of COS cells with NGF (Chittka and Chao 1999). SC1 acts as a transcriptional repressor and may interact with histone deacetylases to repress the promitotic genes cyclin E and B, thereby inducing growth arrest (Chittka et al. 2004). The NRAGE (neurotrophinreceptor-interacting MAGE homolog) protein also interacts with the p75<sup>NTR</sup>, and it is thought to compete with TrkA receptors for binding to a juxtamembrane domain of the receptor (Salehi et al. 2000). Interaction of NRAGE with the p75<sup>NTR</sup> is mediated by a conserved sequence called MAGE homology domain, which also determines the interaction of necdin and MAGE-H1 (two MAGE proteins) with  $p75^{NTR}$ . The interaction of these two MAGE proteins with the  $p75^{NTR}$  is enhanced by ligand stimulation, which may contribute to the recruitment of a signaling complex to induce differentiation (Tcherpakov et al. 2002). Similarly, the p75<sup>NTR</sup> binding partner NRAGE was shown to facilitate cell cycle exit in HEK293 cells (Salehi et al. 2000). Sall2 also interacts and sequesters p75<sup>NTR</sup>, and activation of TrkA receptors is required to induce the nuclear translocation of the protein with consequent activation of target genes. The induction of p21<sup>WAF1/CIP1</sup> may mediate the effects of Sall2 on NGF-dependent cell cycle arrest and growth inhibition (Pincheira et al. 2009). In contrast with SC1, MAGE proteins, and Sall2, which induce cell cycle arrest, Bex1 (brain-expressed X-linked 1) competes with the receptor interacting protein 2 (RIP2) for binding to the p75<sup>NTR</sup>, thereby maintaining proliferation and preventing differentiation. Studies performed in astrocytes, which express p75<sup>NTR</sup> but not TrkA receptors, showed that NGF attenuates the EGF-induced upregulation of cyclins D1 and E, inhibits the interaction between cyclin D1 and Cdk4, which is required for progression from G1 into S phase, and upregulates cyclin kinase inhibitory proteins (Cragnolini et al. 2012). These evidences indicate that p75<sup>NTR</sup> receptors may act at different levels to induce cell cycle arrest in a given cell.

#### 6 Synaptic Plasticity

Neurotrophins were found to modulate synaptic transmission and plasticity. Valuable insight came from observing the peripheral nervous system, particularly the heart sympathetic innervation. This system can be reproduced in vitro by coculturing SGCs neurons with cardiac myocytes where acute application of norepinephrine increases the contraction frequency of myocytes while acetylcholine has the opposite effect (Furshpan et al. 1976). Neurotrophins regulate sympathetic neuron cotransmission by modulating the activity-dependent release of norepinephrine and acetylcholine (Lockhart et al. 1997; Yang et al. 2002). Similarly to other neuronal functions described throughout this chapter, Trk receptors and p75<sup>NTR</sup> function antagonistically. NGF induces potentiation of excitatory noradrenergic transmission that is mediated by the TrkA receptor (Lockhart et al. 1997), while BDNF induces a shift from excitatory to inhibitory (cholinergic) transmission acting through the p75<sup>NTR</sup> (Yang et al. 2002). CaMKII acts downstream of p75<sup>NTR</sup> to control the shift from excitatory to inhibitory transmission since presynaptic inhibition of CaMKII activity or expression of constitutively active CaMKII blocks or mimics BDNF-induced inhibitory transmission respectively (Slonimsky et al. 2006).

Synaptic plasticity in the CNS induced by BDNF has been particularly well studied (Carvalho et al. 2008; Santos et al. 2010). Application of BDNF or NT-3 induced an enhancement of synaptic strength at the Schaffer collateral-CA1 synapses (Kang and Schuman 1995). Accordingly, BDNF regulates long-term potentiation (LTP) in the hippocampus (Figurov et al. 1996; Korte et al. 1996; Patterson et al. 1996) and in the visual cortex (Akaneya et al. 1997; Huber et al. 1998). Application of exogenous BDNF is able to induce LTP in hippocampal slices, and this effect is blocked by BDNF scavengers like BDNF antibodies or TrkB-IgG fusion proteins (Chen et al. 1999; Figurov et al. 1996; Pozzo-Miller et al. 1999). In line with these evidences, BDNF knockout animals show impaired LTP in the CA1 region of the hippocampus, and exogenous BDNF or re-expression of BDNF using a virus-based strategy restored LTP in BDNF-deficient mice (Korte et al. 1996; Patterson et al. 1996). An additional layer of complexity was added when Pang et al. showed tPA is responsible for plasmin-dependent conversion of proBDNF into mature BDNF and this is required for late-phase long-term potentiation (L-LTP) (Pang et al. 2004). However, another study reports proBDNF is rapidly converted intracellularly to mature BDNF and the mature but not the precursor form is secreted upon stimulation. Moreover, the authors show hippocampal long-term depression (LTD) is unaffected in slices prepared from conditional BDNF KO animals (Matsumoto et al. 2008). The reasons for these conflicting results are still elusive and additional studies are required to clarify these divergent observations.

The opposing roles of p75<sup>NTR</sup> and Trk receptors in the nervous system led the investigators to ask if the p75<sup>NTR</sup> could negatively regulate structural and functional plasticity. As hypothesized,  $p75^{NTR-/-}$  mice have higher spine density and greater dendritic complexity than wild-type mice. Accordingly, overexpression of p75<sup>NTR</sup> decreased dendritic complexity and the number of spines (Zagrebelsky et al. 2005). Interestingly the phenotype of p75 (ExonIV-/-) animals, which lack both the FL-p75 and s-p75 isoforms, is more dramatic than the one observed in p75 (EXONIII -/-) animals. The authors claim this could be due to a compensatory effect by the s-p75 isoform (Zagrebelsky et al. 2005). Additionally, Korte and colleagues showed that p75<sup>NTR</sup>-null animals have impaired hippocampal LTD while LTP is unchanged, with the expression levels of GluA2 and GluA3 subunits of the alpha-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) receptor in the hippocampus being altered in the p75<sup>NTR</sup> null animals (Rosch et al. 2005). In line with these studies, proBNDF was shown to induce NMDA-dependent LTD in the hippocampus which is mediated by p75<sup>NTR</sup> and the GluN2B subunit. Blocking the function of p75<sup>NTR</sup> or GluN2B abolishes proBDNF enhancement of LTD (Woo et al. 2005). Furthermore, GluN2B-mediated synaptic currents and expression are reduced in the hippocampus of p75<sup>NTR-/-</sup> mice (Woo et al. 2005). However, at the moment, there is no direct link between p75<sup>NTR</sup> and NMDA receptors and it remains to be determined if p75<sup>NTR</sup> regulates the trafficking or endocytosis of GluN2B and/or of GluA2/3. In Xenopus neuromuscular synapses, exogenous proBDNF induces synaptic depression. Moreover, stimulation-induced release of proBDNF from myocytes leads to synaptic depression and retraction and this effect is mediated by presynaptic p75<sup>NTR</sup> (Yang et al. 2009).

Paralleling the effects attributed to p75<sup>NTR</sup> in synaptic plasticity has proven more difficult in behavioral animal models. Some studies claim p75<sup>NTR</sup> null animals have behavioral impairments, particularly deficits in memory task acquisition (Peterson et al. 1999; Wright et al. 2004), while others presented evidences for better spatial learning performance in p75<sup>NTR</sup> knockout mice when compared to control animals (Barrett et al. 2010; Greferath et al. 2000). The different genetic background of the animals used in these studies could account for the conflicting results, but further investigation is required to reconcile this discrepancies. Taken together, these results show that p75<sup>NTR</sup> has a role in activity-dependent synaptic plasticity.

## 7 Conclusion

Although p75<sup>NTR</sup> was the first neurotrophin receptor to be identified, for many years, it stood in the shadow of the Trk receptor with a mere supporting role. We now know that p75<sup>NTR</sup> plays a major role in the nervous system, especially during development and in injury scenarios, but with the latest developments, we face the

risk of this vision also becoming outdated. Recent studies provided a fresh new look into p75<sup>NTR</sup> function, for example, the diversity of mechanisms induced to promote cell cvcle arrest in a given cell, its role in axonal pruning, and the specificity of neural connectivity in the adult brain (Park et al. 2010; Singh et al. 2008). Also, unforeseen p75<sup>NTR</sup> partners and consequently new functions were identified, adding new layers of complexity to p75<sup>NTR</sup> function. It will be of the outmost importance to design experiments to understand how these different ligands, coreceptors, and cytosolic interactors, acting through the same receptor, activate the specific signaling pathways that are responsible for the variety of physiological functions attributed to the  $p75^{NTR}$ . It is tempting to speculate that the different microdomains within a neuronal cell could provide the most needed functional specificity but this is unlikely to be the sole answer to this question. A major challenge will be to understand the spatial and temporal expression patterns of p75<sup>NTR</sup>, its co-receptors, homologues, and intracellular partners, to have an integrative view of  $p75^{NTR}$  function in the nervous system.

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# p75NTR: A Molecule with Multiple Functions in Amyloid-Beta Metabolism and Neurotoxicity

Yan-Jiang Wang, Fan Zeng, Khalil Saadipour, Jian-Jun Lu, and Xin-Fu Zhou

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Y.-J. Wang (⊠) • F. Zeng Department of Neurology, Daping Hospital, Third Military Medical University, Daping, Chongqing, China e-mail: yanjiang\_wang@tmmu.edu.cn; qwiwp@126.com

K. Saadipour • J.-J. Lu

Division of Health Sciences, University of South Australia, Adelaide, SA, Australia e-mail: khsaadipour@gmail.com; stevelu90@hotmail.com

#### X.-F. Zhou

School of Pharmacy and Medical Sciences, Faculty of Health Sciences, University of South Australia, Adelaide, SA, Australia e-mail: xin-fu.zhou@unisa.edu.au

#### Abstract

The p75 neurotrophin receptor (p75NTR) is a pan-receptor for neurotrophins including nerve growth factor, brain-derived neurotrophic factor, neurotrophin-3, and neurotrophin 4/5. P75NTR plays a diverse role from regulating cell survival to modulating neurite outgrowth. Under some pathological conditions, p75NTR expression is activated and plays detrimental roles in disease progression. Alzheimer's disease (AD), the most common form of dementia, is characterized by the deposition of amyloid plaques, accumulation of fibrillary tangles in neurons, neurite degeneration, loss of neurons, and a progressive loss of cognitive function. Recent studies suggest that p75NTR, also a receptor for amyloid-beta (A $\beta$ ), is a critical factor involved in the pathogenesis of AD. This chapter is to discuss the roles of p75NTR in the production of amyloidbeta  $(A\beta)$ , neuronal death, neurite degeneration, tau hyperphosphorylation, cell cycle re-entry, and cognition decline, to propose that p75NTR is a potential target for the development of therapeutic drugs for AD, and to provide perspectives in developing various therapeutic strategies targeting different aspects of AD hallmarks which relate to p75NTR functions, and breaking the p75NTR-mediated positive-feedback loop which promotes the cascades in the pathogenesis of AD.

#### Keywords

Alzheimer's disease • Amyloid-beta • Cell cycle • Cognition • Extracellular domain • Neurodegeneration • Neuronal death • p75NTR • tau

List of Abbreviations				
AD	Alzheimer's disease			
APP	Amyloid precursor protein			
Αβ	Amyloid-beta			
BACE	Beta-site APP-cleaving enzyme			
BDNF	Brain-derived neurotrophic factor			
CNS	Central nervous system			
DRG	Dorsal root ganglion			
ECD	Extracellular domain			
MAG	Myelin-associated glycoprotein			
NBM	Nucleus basalis of Meynert			
NGF	Nerve growth factor			
NT	Neurotrophin			
p75NTR	p75NTR neurotrophin receptor			
PNS	Peripheral nervous system			
TACE	TNF-alpha converting enzyme			
TNF	Tumor necrosis factor			
Trk	Tropomyosin receptor kinase			
Aβ BACE BDNF CNS DRG ECD MAG NBM NGF NT p75NTR PNS TACE TNF	Amyloid-beta Beta-site APP-cleaving enzyme Brain-derived neurotrophic factor Central nervous system Dorsal root ganglion Extracellular domain Myelin-associated glycoprotein Nucleus basalis of Meynert Nerve growth factor Neurotrophin p75NTR neurotrophin receptor Peripheral nervous system TNF-alpha converting enzyme Tumor necrosis factor			

## 1 Introduction

The p75 neurotrophin receptor (p75NTR) was first identified about 20 years ago as the receptor for nerve growth factor (NGF) and was subsequently found to be the receptor for all other neurotrophins including brain-derived neurotrophic factor (BDNF), neurotrophin-3 (NT-3), and neurotrophin-4/neurotrophin-5 (NT-4/NT-5) (Chao and Bothwell 2002; Barker 2004). p75NTR belongs to the tumor necrosis factor receptor superfamily which consists of ten different transmembrane glycoproteins (type I) (Gruss 1996). The role as the pan-receptor for neurotrophins enables p75NTR to play a diverse role in regulating cell survival. It can activate at least three signaling pathways including neuronal growth, apoptosis, and synapse plasticity, and most recently this receptor is also found to be involved in the neuronal cell cycle re-entry, which is considered a key pathological mechanism underlying the development of Alzheimer's disease and other neuronal degenerative diseases. This chapter focuses on the recent progress in understanding the physiological and pathological roles of p75NTR with particular interest in the role of p75NTR in development and intervention of Alzheimer's disease.

## 2 Biology of p75NTR

## 2.1 Molecular Biology, Structure, Distribution, and Expression Pattern

The structure of the human p75NTR gene was first described by Santee et al. in 1996 (Santee and Owen-Schaub 1996). It was identified as 19728 base pairs (bp) in size containing six exons and five introns. Sharing an analogous structure and function with the tumor necrosis factor (TNF) receptor, p75NTR is a transmembrane protein that consists of an extracellular domain, a transmembrane domain, and an intracellular domain. The extracellular domain consists of a stalk domain which links the transmembrane domain and four cysteine-rich repeat domains that facilitate neurotrophin binding. There is clear evidence that the third and fourth cysteine repeats are the binding sites for neurotrophins, as the mutant form of p75NTR that lacks these two domains cannot bind neurotrophins. The intracellular part of p75NTR contains an 80-amino acid "death domain" module. It connects the transmembrane domain through a flexible linker region N-terminal domain. The flexibility of the linker domain possibly plays an important role in multiple signal transduction. In addition, the intracellular, cytoplasmic region of p75NTR does not exhibit an intrinsic ligand-inducible enzymatic function, and this is distinctly different from tropomyosin receptor kinase (Trk) receptors. Although p75NTR is a member of the TNF receptor family, there are some differences between p75NTR and other TNF receptors. The death domain of p75NTR is a type-2 molecule, which is different from the type-1 molecules of other TNF receptors. Furthermore, unlike other TNF receptors, p75NTR does not self-associate in solution.

## 2.2 Expression Location of p75NTR

The p75NTR has a very wide tissue and cellular distribution in a wide range of species and its distribution varies during the different stages of the life. Its expression has been shown at the early developmental stages of the central nervous system (CNS) and peripheral nervous system (PNS). In the developing rat dorsal root ganglion (DRG), p75NTR expression can be found as early as E7 (Ernfors et al. 1988). The receptor was also found to be selectively expressed in the sympathetic ganglia (Buck et al. 1987). In the basal forebrain of the rat, the p75NTR immunoreactivity can be observed as early as E15 (Koh and Loy 1989) and throughout the whole of the adult life (Mufson and Kordower 1992; Hartig et al. 1998). In contrast to the other tissue organs, the expression of p75NTR in these basal forebrain neurons seems to be upregulated during aging while the Trk expression is downregulated (Ginsberg et al. 2006). Interestingly, p75NTR levels tend to be high and widely expressed during development but decrease dramatically in adulthood. In addition, p75NTR can be reexpressed or upregulated to a significant degree in pathological states related to neural cell death or after injury (Barrett and Bartlett 1994). In the adult mouse, p75NTR expression has also been shown to be present in a small subset of spinal cord ascending sensory axons, and in some retinal cells and subsets of dorsal root ganglia, but not the cortex and cerebellum. In addition to neurons, p75NTR is also expressed in a variety of glial tissues such as Schwann cells and oligodendrocytes during development and after injury (Cragnolini and Friedman 2008).

#### 2.3 Physiological Roles of p75NTR and Its Metabolism

In contrast to the Trks receptors, the physiological roles of p75NTR are complicated and difficult to discern. Analysis of the p75NTR function has been complicated by the fact that the major physiological role of p75NTR may change dramatically depending on the cell context. p75NTR is a low-affinity pan-receptor which can functionally collaborate with Trk receptors to either enhance or reduce neurotrophin-mediated Trk receptor activation. The binding of p75NTR and Trks to neurotrophins can trigger several signal transduction pathways, which subsequently promote the survival, differentiation, and development of neuronal cells. In addition to promoting survival of neuronal tissues, another well-established and important functional role of p75NTR is to mediate the apoptosis in normal and pathological conditions. p75NTR has been shown to induce apoptosis in different cell types, including cultured neonatal sympathetic neurons (Bamji et al. 1998), motor neurons (Sedel et al. 1999), sensory neurons (Barrett and Bartlett 1994), oligodendrocytes (Casaccia-Bonnefil et al. 1996), and embryonic proprioceptive neurons when Trk activation is reduced or absent (Davey and Davies 1998). There is evidence indicating that p75NTR is involved in cholinergic neuronal death independent of TrkA expression during development and after injury (Van der Zee et al. 1996; Naumann et al. 2002).

p75NTR has more recently been shown to regulate neurite outgrowth through regulation of RhoA, in some contexts following activation by myelin proteins in conjunction with the Nogo66 receptor (Barker 2004). It can also promote neuronal differentiation and cell cycle. Adult dorsal root ganglion neurons or postnatal cerebellar neurons from mice carrying a mutation in the p75NTR gene are insensitive to MAG with regard to neurite collapse (Yamashita et al. 2002).

Some novel functional roles of p75NTR have been identified recently. NGF stimulates neurite outgrowth from embryonic rat hippocampal neurons and chick ciliary neurons which express p75NTR but not TrkA (Brann et al. 1999). The p75NTR receptor may also be involved in the inhibition of axonal elongation by myelin. In mice lacking p75NTR, aberrant axonal elongation is observed in the myelin-rich area where these axons would not grow (Walsh et al. 1999). These findings suggested that p75NTR may have bidirectional roles in regulating neurite outgrowth (Hasegawa et al. 2004).

In a recent study, a novel p75NTR-interacting protein named Sall2 has been found to link NGF signaling to cell cycle progression and neurite outgrowth (Pincheira et al. 2009).

p75NTR is physiologically cleaved by the enzyme known as TNF-alpha converting enzyme (TACE) in the juxtamembrane domain, to shed the extracellular domain (Weskamp et al. 2004). The exact significance of this process remains unknown. It is possible that the shedded extracellular domain can play an opposing role to negatively regulate the functions of the membrane counterpart, by competitively inhibiting the binding of ligands to membrane p75NTR.

#### 2.4 Signal Pathway of p75NTR Action

p75NTR has a close collaboration with tyrosine kinases receptors. It regulates ligand specificity of Trk receptors and modulates binding affinity of TrkA to NGF (Huang and Reichardt 2003). Although other neurotrophins can interact with TrkB receptors, only BDNF provides a functional response upon co-expression of TrkB with p75NTR. Similarly, both NGF and NT-3 bind to TrkA, but p75NTR restricts TrkA-NGF signaling. In contrast, co-expression of p75NTR and TrkC decreases ligand specificity of TrkC to NT-3.

p75NTR interacts with a dozen of signaling molecules and mediates apoptosis, Schwann cell migration, myelination, axonal growth, and regeneration. The intracellular domain of p75NTR contains several potential motifs which modulates interactions of p75NTR with other proteins, subsequently activating different signaling pathways (Barker 2004; Bronfman and Fainzilber 2004; Dechant and Barde 2002; Roux and Barker 2002) such as NF-κB (Carter et al. 1996), c-jun kinase (JNK) (Friedman 2000; Yoon et al. 1998), and caspases (Troy et al. 2002). In addition, p75NTR activates a small GTPase Rho (Yamashita et al. 1999) involving different ligands derived from myelin, such as myelin-associated glycoprotein (MAG) and Nogo (Wang et al. 2002a; Wong et al. 2002; Yamashita et al. 2002) and two different co-receptors: a lipid-anchored ligand-binding subunit known as Nogo receptor (NgR) (Fournier et al. 2001) and Lingo-1 (Mi et al. 2004). For instance, BDNF binds to p75NTR and induces apoptosis through activation of JNK pathway (Bhakar et al. 2003; Palmada et al. 2002). p75NTR and sortilin form a very-high-affinity receptor complex for the binding of proNGF and other proneurotrophins which promote apoptosis (Nykjaer et al. 2004). p75NTR also mediates biological activity of MAG and Nogo which inhibit axonal elongation and neurite growth via interacting with the Nogo receptor (Wang et al. 2002a, b).

## 3 Pathological Roles of p75NTR in the Nervous System

Although the expression of p75NTR is developmentally regulated in the nervous system, marked increase is observed under certain pathological conditions; however, the significance of the expression in these conditions is still not very clear.

In the dorsal root ganglia, the reduction of p75NTR levels by antisense oligonucleotides prevents the loss of axotomized neurons (Cheema et al. 1996). Similarly, the loss of motor neuron occurring after transection of the neonatal facial nerve is attenuated in mice carrying a mutation in the p75NTR gene (Ferri et al. 1998; Ferri and Bisby 1999). In contrast, the administration of NGF into the transected neonatal facial nerve of animals increases cell death (Sendtner et al. 1992). p75NTR is reexpressed in the transected neurons at levels comparable to those seen during the developmental stages. The upregulation of p75NTR is also observed in the cerebral cortex in Alzheimer's disease (Mufson and Kordower 1992). In a mouse model of amyotrophic lateral sclerosis, in which a mutant form of superoxide dismutase is overexpressed, p75NTR is reexpressed in lumbar motor neurons (Lowry et al. 2001) as the disease progresses. Consistently, the expression of p75NTR is also observed in motor neurons in the cervical spinal cords of patients with amyotrophic lateral sclerosis.

After axotomy, cortical spinal neurons strongly reexpress p75NTR three days after lesion, when neuronal death occurs. The activation of p75NTR by NT-3 causes the death of these neurons, as shown by experiments using blocking antibodies against p75NTR or NT-3 (Giehl et al. 2001). In spinal cord injury, p75NTR is essential for neuronal cell survival and functional recovery after injury (Chu et al. 2007). Activation of p75NTR contributes to NMDAR-mediated apoptosis in the neonatal brain (Griesmaier et al. 2010). These findings further suggest that p75NTR may be involved in neuronal death in neurological disorders. In a study using experimental autoimmune encephalomyelitis (EAE) animal models, in contrast to the clinical manifestation of EAE observed in wild-type C57/BL6 mice, mice deficient of p75NTR developed severe or lethal diseases and concomitant increased levels of inflammation in the CNS (Copray et al. 2004). This may suggest p75NTR is involved in the inflammation process but the detailed molecular mechanism is not clear. In retina degeneration, p75NTR is upregulated and exerts neurotoxic effect through paracrine mechanism (Bai et al. 2010). It appears that in pathologies involved in neuronal degeneration and death, p75NTR expression has a tendency to be upregulated in most experimental models used so far. However, the mechanism underlying the upregulation is not completely understood.

## 4 Roles of p75NTR in Pathogenesis of Alzheimer's Disease

#### 4.1 p75NTR Expression in the Brain of Alzheimer's Disease

It is well known that cholinergic neurons in the basal forebrain especially in the nucleus basalis of Meynert (NBM) are most vulnerable and severely affected in Alzheimer's disease and they express p75NTR-positive, TrkA, TrkB, and TrkC (Salehi et al. 2000). Strikingly, very little or no p75NTR-positive neurons are normally observed in other brain regions. How the level of p75NTR expression changes in Alzheimer's disease patients and during the aging process remains controversial. Some researchers find that the expression of the nerve growth factor (NGF) receptor (including p75NTR) is increased in both cortical neurons and CA1 and CA2 subfields of the hippocampus in Alzheimer's disease patients compared with controls (Mufson and Kordower 1992; Hu et al. 2002). Some other studies show a significant reduction of p75NTR in Alzheimer's disease brains (Arendt et al. 1997; Salehi et al. 2000). Meanwhile, no significant differences in expression levels for p75NTR were observed across control, mild cognitive impairment (MCI, a pre-stage of Alzheimer's disease), nor Alzheimer's disease in other reports (Goedert et al. 1989; Treanor et al. 1991; Mufson et al. 2002; Ginsberg et al. 2006). The discrepancy on p75NTR expression in Alzheimer's disease may be due to the differences in investigated brain regions and measurements. The severity of Alzheimer's disease also influences the level of p75NTR, because it is certain that the greater the loss of cholinergic neurons occurring in Alzheimer's disease, the less that p75NTR is present in the basal forebrain. More work is required before a definite conclusion about this problem can be drawn regarding this issue. Studies on p75NTR expression in the Alzheimer's disease animal models and cell lines provide a clearer answer. In two different strains of transgenic Alzheimer's disease mice, the triple transgenic mice (3xTg-AD, harboring PS1M146V, APPswe, and tauP301L) and APPswe/PS1dE9 mice, the level of p75NTR is significantly increased and associated with the age-dependent rise of A $\beta$ . Meanwhile, A $\beta$  has promoted p75NTR expression in the membrane of SH-SY5Y human neuroblastoma cells (Chakravarthy et al. 2010). They further demonstrated that the microinjected  $A\beta$  oligomer can also increase p75NTR expression, which according to the results of in vitro study may be mediated through an insulin-like growth factor 1 receptor (IGF-1R) phosphorylation (Ito et al. 2012). Consistently, it was found that p75NTR expression increases in an agedependent manner in wild-type mice and is higher in APPswe/PS1dE9 mice relative to their wild-type littermates at the same age (Wang et al. 2011). Although many of the human studies show a reduction of the p75NTR level in the brain, reflecting the loss of p75NTR-positive cholinergic neurons, these animal studies suggest that p75NTR expression increases with aging and is further activated by  $A\beta$ . The neurons of the transgenic animal models are more resistant to  $A\beta$  toxicity as reflected by the loss of very few neurons in the brain of the animal models, relative to humans (Gotz and Ittner 2008). Moreover, it was recently discovered that p75NTR would express in the cultured epithelial cells of choroid plexus, mainly in the cytoplasm and CSF side (Spuch and Carro 2011), which is an indication of its potential role to transport  $A\beta$ through BBB (Zhou and Wang 2011).

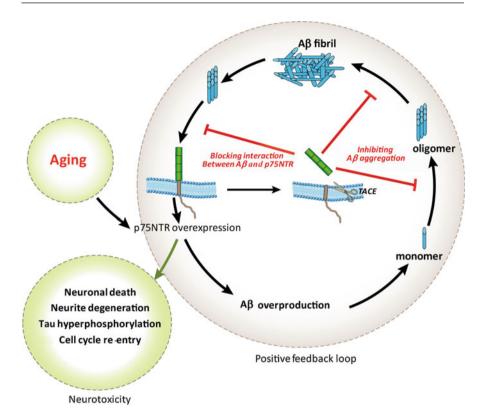
## 4.2 p75NTR Pathway Promotes Aβ Production

The level of p75NTR is significantly elevated in Alzheimer's disease brains, but little is known about the reason and meaning of this change. In our studies, the expression of p75NTR is much higher in the neurons of APPswe transgenic mice than that of the wild-type, suggesting A $\beta$  is very likely to cause the high level of p75NTR in Alzheimer's disease. Our further studies showed that after the p75NTR gene knockout in APPswe transgenic mice, the production of A $\beta$  was reduced in the cortex neurons, and the level of soluble A $\beta$  also decreased (Wang, Wang et al. 2011). This indicates that p75NTR is able to promote A $\beta$  production. Based on these results, it is speculated that A $\beta$  upregulates the expression of p75NTR in Alzheimer's disease brains, and the activated p75NTR in turn stimulates A $\beta$  production. Consequently they form a vicious cycle and finally result in A $\beta$  overproduction. This hypothesis can explain the reason why A $\beta$  production accelerates in Alzheimer's disease brains (Kawarabayashi et al. 2001) and why sporadic Alzheimer's disease patients have overproduction of A $\beta$ when they do not have any known gene mutations with A $\beta$  production (Selkoe 2001). It is demonstrated that beta-site APP-cleaving enzyme 1 (BACE1) protein increases in an annulus around amyloid plaques cores, and it occurs without any change in BACE1 mRNA level. Therefore, it can be concluded that amyloid plaques lead to the increased level of BACE1 surrounding neurons at an early stage before neuronal death occurs, and this may drive a positive-feedback loop in A $\beta$  production (Zhao et al. 2007) (Fig. 1). Consistent with this conclusion, further research has described the same interactions between BACE1 and  $A\beta$  in the pathogenesis of Alzheimer's disease, no matter whether it is familial or sporadic (Tabaton et al. 2010). The positive-feedback loop in A $\beta$  production is an important issue worthy of further investigations (Fig. 1).

Aging is the most important risk factor for Alzheimer's disease. Recently some researchers have identified the p75NTR system as a molecular link between normal aging and Alzheimer's disease. When p75NTR, which activates  $\beta$  cleavage of APP, is upregulated and TrkA, which reduces the cleavage, is downregulated during aging, the p75NTR-ceramide pathway is significantly activated, causing the increase of steady-state levels and activity of BACE1 downstream and eventually resulting in the acceleration of A $\beta$  production (Costantini et al. 2005b). The caloric restriction and nSMase inhibitors could reduce A $\beta$  production and the risks associated with aging, by interrupting this signaling pathway. Both the upregulation of p75NTR expression and the acceleration of A $\beta$  production follow an age-dependent fashion. Further study of this research demonstrated that signaling through IGF1-R controls the level of TrkA and p75NTR in both human neuroblastoma cell lines and primary neurons from the mouse brain, and IGF1-R regulates A $\beta$  production in the upstream of p75NTR-ceramide pathway (Costantini et al. 2006).

### 4.3 p75NTR Regulates Aβ Deposition and Clearance

 $A\beta$  mainly deposits in the cerebral neocortex, hippocampus, and vessel walls, which are coincident with the projection area of cholinergic neurons and



**Fig. 1** Schematic diagram of p75NTR functions in Alzheimer's disease. In the brain of Alzheimer's disease, p75NTR expression is upregulated by aging and further activated by  $A\beta$ . p75NTR signaling increases  $A\beta$  production and thus enhances steady-state levels of  $A\beta$  in the interstitial space which may increase AD pathology.  $A\beta$  again promotes p75NTR expression. P75NTR/A $\beta$  forms a positive-feedback loop. Besides  $A\beta$  production, p75NTR also mediates neurotoxicity of  $A\beta$ , such as neuron death, neurite degeneration, tau hyperphosphorylation, and cell cycle re-entry. p75NTR is to be cleaved by an enzyme named TACE to shed its extracellular domain. The extracellular domain of p75NTR after shedding from the membrane may bind to the 25–35 amino acid of  $A\beta$ , which is mainly responsible to  $A\beta$  aggregation and neurotoxicity, thus suppressing  $A\beta$  aggregation and reducing  $A\beta$  deposition in the brain. Meanwhile, the extracellular domain of p75NTR may also block the interaction between  $A\beta$  and membrane p75NTR by competitive binding  $A\beta$ , thus attenuating the p75NTR signaling which leads to both  $A\beta$  production and neurotoxicity. The shedded extracellular domain of p75NTR might be an endogenous protective molecule against  $A\beta$  production, deposition, and toxicity and would be an important and promising therapeutic target of Alzheimer's disease

sympathetic neurons expressing p75NTR. The underlying mechanism remains unclear. It was found that in the cerebral cortex and hippocampus of APPswe transgenic Alzheimer's disease mice, more than 90 % of the A $\beta$  plaques contained p75NTR-positive fibers. More importantly, the p75NTR-positive fibers are mostly located in the cores of A $\beta$  plaques, while the p75NTR-negative degenerated ones are located around the plaques. The appearance of p75NTR-positive neurites is prior to A $\beta$  plaques in the brain of Alzheimer's disease mice. These findings all indicate that p75NTR expressed on the fibers is probably an initiating or promoting factor for A $\beta$  deposition. However, deletion of p75NTR can increase total A $\beta$  level in the brain, with increased fibrillar A $\beta$  levels but decreased soluble A $\beta$  level. Therefore, how p75NTR influences A $\beta$  deposition may be complicated.

As a member of the tumor necrosis factor receptor superfamily, p75NTR is cleaved by metalloproteases mainly TACE or ADAM17 in the normal metabolic process, and then the extracellular domain of p75NTR (p75NTR-ECD) is released (Weskamp et al. 2004). The physiological function of p75NTR-ECD has not been well studied. In our studies, after deleting the p75NTR gene of APPswe transgenic mice, insoluble  $A\beta$  deposition is significantly increased in the brain (Wang, Wang et al. 2011). Further study demonstrates that recombinant p75NTR-ECD inhibits A $\beta$  oligometization in a dose-dependent manner in vitro, and injection of p75NTR-ECD in the hippocampus of Alzheimer's disease mice can reduce local Aß plaques. Based on these results, it can be concluded that p75NTR-ECD, which is released during the metabolic process of p75NTR, is a key regulatory factor of inhibiting A $\beta$  aggregation and facilitating disaggregation of A $\beta$  fibrils. It may act as an A $\beta$ -sequestering molecule to clear A $\beta$  in vivo. Seeking the A $\beta$  binding sites on p75NTR-ECD is very valuable for revealing the mechanism about how p75NTR regulates A $\beta$  deposition. After that, the effective control and interference with the deposition and clearance of  $A\beta$  can be achieved. It is worthy to note that since p75NTR is also expressed in endothelial cells of BBB, it may help transport A $\beta$ from CSF to blood, thus facilitating A<sup>β</sup> clearance to some extent (Zhou and Wang 2011).

In the progression of Alzheimer's disease, the level of p75NTR is related to  $A\beta$  load and other factors in the brain. How is the extracellular shedding of p75NTR regulated in vivo? Is there any change of p75NTR-ECD level in the Alzheimer's disease brain, cerebral spinal fluid, blood, or urine? Are the function and metabolism of p75NTR-ECD impaired in Alzheimer's disease patients? Can  $A\beta$  influence p75NTR-ECD production? By what means does p75NTR transport  $A\beta$  through BBB? Are there any single nucleotide polymorphisms of the p75NTR gene associated with the risk of Alzheimer's disease? All these important questions need to be answered in future studies.

### 4.4 p75NTR and Neurite Degeneration

Neurite degeneration is a pivotal event of Alzheimer's disease and a main contributor of the memory loss. A $\beta$  is an important factor which leads to neurite degeneration in Alzheimer's disease patients, but its receptor pathway and mechanism are still unknown (Petratos et al. 2008). It is demonstrated that intact p75NTR mediates or enables A $\beta$ -induced neurite degeneration in vitro and in vivo via the c-Jun pathways, a known mediator of A $\beta$ -induced deleterious signaling. After the removal of p75NTR extracellular domain, the A $\beta$ -induced neuritic dystrophy in cultured hippocampus neurons and in the basal forebrain of chronic Alzheimer's

disease mice could be completely reversed, and the neuritic dystrophy in the hippocampus was also significantly diminished (Knowles et al. 2009). Besides, non-peptide small molecule p75NTR ligands which block the binding of A $\beta$  to p75NTR are found to be capable of inhibiting A $\beta$ -induced neurite degeneration, synapse impairment, and the activation of some molecules involved in Alzheimer's disease pathology including calpain/cdk5, GSK3β and c-Jun, and tau phosphorylation. They can also prevent A $\beta$ -induced inactivation of AKT and CREB (Yang et al. 2008). In accordance with results above, it was observed that abundant degenerative fibers and neurites in the neocortex and hippocampus express p75NTR in APPswe/PS1dE9 mice (Wang et al. 2010b), suggesting that p75NTR is a key receptor modulating A $\beta$ -induced neurite degeneration. At the same time, a recent study found that the sympathetic nervous system is severely impaired in p75NTR-deficient Alzheimer's disease model, which leads to critical dysfunction of sympathetic innervation to multiple organs and finally results in the death of the animal. The protective role of p75NTR against A $\beta$ -induced neurite degeneration showed in this study could be the result of the combined effect of p75NTR and its co-receptors under in vivo condition, such as the enhanced NGF and TrkA signaling, and possibly the promoted glial cell line-derived neurotrophic factor (GDNF) signaling, especially in the sympathetic nervous system (Bengoechea et al. 2009).

The reduction of NGF has also been suggested to be involved in the pathogenesis of Alzheimer's disease, but the exact role of its precursor (proNGF) in Alzheimer's disease remains elusive. It is reported that proNGF is the predominant form of this neurotrophin in the human brain, and it binds to p75NTR more strongly than mature NGF (Fahnestock et al. 2001). ProNGF has been shown to induce apoptosis in neuronal cell cultures through its interaction with p75NTR and its co-receptor, sortilin (Niewiadomska et al. 2011). Besides, the binding of proNGF to p75NTR-SorCS2 (sortilin-related VPS10 domain-containing receptor 2) complex can disassociate the guanine nucleotide exchange factor Trio from the complex, which brings down the activity of the small GTPase Rac and finally leads to destabilization and collapse of actin filaments, as well as growth cone retraction and collapse in cultured hippocampus neurons (Deinhardt et al. 2011). Since the level of proNGF has been found to increase in the cerebral cortex in Alzheimer's disease, it may mediate neurite degeneration through this mechanism. Besides, blocking proNGF can inhibit neuronal loss (Friedman 2000; Podlesniy et al. 2006; Volosin et al. 2008). The effects of proNGF in vitro and in vivo were investigated, and the results showed that proNGF is localized to the A $\beta$  plaques in the brain of Alzheimer's disease mice, which also induced the neuronal death and neurite degeneration of different neuronal cell lines and primary neurons in culture, likely dependent on the expression of p75NTR (Wang et al. 2010a) (Fig. 1).

### 4.5 p75NTR and Neuronal Death

Neuronal death is another important pathological character of Alzheimer's disease. As one ligand of p75NTR,  $A\beta$  is best known for mediating neuronal death and has

been consistently linked to the pathology of Alzheimer's disease. It is believed that the interactions of A $\beta$  with p75NTR and other neuronal cell surface receptors can trigger the production of reactive oxygen species, expression of caspases and pro-apoptotic genes (such as p53, p35), and enhance mitochondrial permeability (Dias Bda et al. 2011). Evidence shows that p75NTR plays a critical role in the early and characteristic loss of cholinergic neurons in response to  $A\beta$  in the septohippocampal pathway in Alzheimer's disease (Sotthibundhu et al. 2008). A recent study demonstrated that overexpression of tumor necrosis factor receptor-associated factor 6 (TRAF6) and p62, which are essential in p75NTR polyubiquitination and NF-kB activation-mediated cell survival signaling downstream, can reduce A $\beta$ -induced neuronal death (Geetha et al. 2012). It was found that in the presence of A $\beta$ , more neurons survived in culture after the p75NTR gene was deleted. Besides, the application of the recombinant extracellular domain of p75NTR can decrease the A $\beta$ -induced neuronal death. Thus, it is probable that A $\beta$ toxicity can be antagonized by blocking the p75NTR-mediated pathway (Wang et al. 2010b). However, research shows that the role of p75NTR in neurotoxicity depends on the different forms of A $\beta$  interacting with it. For fibrillar A $\beta$ , p75NTR mediates neuronal death, whereas for soluble oligometric A $\beta$ , it exerts a protective role against the neurotoxicity (Costantini et al. 2005a). The protective effect may be mediated by phosphatidylinositide 3-kinase (PI3K) activity due to the function of the juxtamembrane sequence of the cytoplasmic region of p75NTR. Considering that the activation of p75NTR-mediated pathway may need the role of its specific co-receptors, and that most studies were confined to in vitro experiments, further investigation about how p75NTR mediates Aβ-induced neurotoxicity needs to be carried out in in vivo models (Coulson 2006).

proNGF participates in neuronal death in Alzheimer's disease as well. Some researchers demonstrate that proNGF from the injured spinal cord induces apoptosis in culture among p75NTR-positive, but not p75NTR-negative, oligodendrocytes, and its action can be blocked by proNGF-specific antibodies. ProNGF can eliminate damaged cells by activating the apoptotic machinery of p75NTR (Beattie et al. 2002). Other researchers also found that proNGF secreted by astrocytes under the stimulation of peroxynitrite can specifically cause motor neuron death via the engagement of p75NTR (Domeniconi et al. 2007). As for Alzheimer's disease, some studies showed that proNGF levels increased in glial cells and neurons of the cortex and hippocampus, which is closely related to the progression of Alzheimer's disease. The upregulation of proNGF may represent the dysfunction of maturation of NGF that enhances the vulnerability of cholinergic neurons in Alzheimer's disease (Cuello et al. 2007). Moreover, researchers found proNGF extracted from Alzheimer's disease brains can obviously induce apoptotic neuronal death through p75NTR (Pedraza et al. 2005). And p75NTR intracellular domain (p75NTR-ICD), the level of which was significantly increased and paralleled the increase of proNGF in the entorhinal cortex, should account for proNGF-induced apoptosis in Alzheimer's disease (Podlesniy et al. 2006). Nevertheless, some researchers speculated the possibility that the loss of proNGF-mediated signaling may reduce the TrkA receptors within the cholinergic basal forebrain (CBF) neurons, affecting cell survival. Whether proNGF binds p75NTR in vivo leading to apoptosis remains unclear (Peng et al. 2004). A recent study indicated that cleavage-resistant mutated forms of proNGF can promote neurite outgrowth, increase cell survival, as well as cause apoptotic cell death through phosphorylation of TrkA and ERK1/2 or initiation of apoptotic pathway by cleavage of caspase 3. So the final result of cell viability or apoptotic effects will depend on the balance of proNGF:NGF and p75NTR:TrkA ratios in Alzheimer's disease brains (Clewes et al. 2008). This is further proved as deletion of p75NTR-ECD which consequently leads to the abrogating of proNGF signaling in AD12 mice, obtained by crossing AD10 mice which express anti-NGF antibodies with p75NTR<sup>exonIII(-/ -)</sup> mice, can still exert a protective role to anti-NGF-induced amyloidogenesis as a result of dominant TrkA survival signaling (Capsoni et al. 2010).

#### 4.6 p75NTR and Neuronal Cell Cycle Re-entry

Adult neurons in the brain are classically thought to be in terminal differentiation permanently, effectively precluding them from cell division. However, around a decade ago, it was shown that susceptible cortical and hippocampal neurons in Alzheimer's disease display an activated cell cycle, suggesting cell cycle re-entry of neurons. Accumulating evidence indicates that neuronal cell cycle re-entry represents an early and critical event in Alzheimer's disease, recapitulating known hallmarks of the disease including NFTs and deposition of A $\beta$  plaques. The cell cycle re-entry hypothesis may be extensively applicable in many neurological diseases because the re-entry of cell cycle also occurs in other neurodegenerative diseases.

The mechanism and the role of p75NTR in cell cycle re-entry in Alzheimer's disease remain obscure. However, there is evidence that illustrates an important position of p75NTR in cell cycle re-entry. Firstly, it was demonstrated that during development of the nervous system, the activation of p75NTR by an endogenous source of NGF (possibly proNGF) can trigger cell cycle re-entry and tetraploidization of retinal ganglion cells (Jansen et al. 2007; Nakamura et al. 2007; Morillo et al. 2010). Secondly, many newly identified p75NTR intracellular interactors have been known to regulate cell cycle. Finally, as a known stress-response receptor, expression of p75NTR in affected neurons in Alzheimer's disease becomes upregulated in response to stress, which has a close relationship with cell cycle re-entry. The exact role of p75NTR in the cell cycle re-entry of neurons in Alzheimer's disease needs further investigation.

p75NTR can be activated by A $\beta$  (Yaar et al. 1997) and promote neuritic dystrophy and apoptosis in vitro and in vivo (Sotthibundhu et al. 2008; Knowles et al. 2009). Binding of aggregated A $\beta$  to p75NTR can also lead to Ras/ERK activation (Susen and Blochl 2005), suggesting that the previously described increase of Ras/ MAPK activity in the Alzheimer's disease brain could be induced by this particular p75NTR-dependent signaling pathway (Arendt et al. 1995; Gartner et al. 1995). Furthermore, p75NTR has also been shown to physically

and functionally interact with APP (Fombonne et al. 2009). p75NTR can interact with different neurotrophin precursors, including proNGF (Fombonne et al. 2009) which represents the predominant form of NGF in the brain. This interaction requires the presence of sortilin, known to bind to the "pro" domain. Interestingly, the expression of sortilin becomes increased in aging basal forebrain neurons (Al-Shawi et al. 2007; Al-Shawi et al. 2008), and this molecule is required for neuronal degeneration associated with aging (Jansen et al. 2007). ProNGF has been shown to trigger pro-apoptotic signals through the p75NTR/sortilin complex (Nykjaer et al. 2004) although proNGF-dependent trophic effects are also observed in vivo (Fahnestock et al. 2004; Al-Shawi et al. 2008). Importantly, proNGF becomes accumulated in the cortex and hippocampus (Fahnestock et al. 2001) of the Alzheimer's disease brain, as well as in the aged brain.

## 4.7 p75NTR and Cognition, Tau Phosphorylation

In Alzheimer's disease patients degenerated cholinergic neurons in the basal forebrain are closely related to the impaired learning and memory processes. Some researchers found that after deletion of exon III or IV of the p75NTR gene which leads to a partial or complete p75NTR knockout, the cholinergic neurons in the medial septal nucleus increase 13 % and 28 %, respectively, in Alzheimer's disease mice, which is sure to improve cognition to some degree (Naumann et al. 2002). The results also suggest that the cytoplasmic domain of p75NTR is responsible for the cell death effect, and complete deletion of the full-length p75NTR can decrease cholinergic neuronal death.

There is also some evidence to suggest that p75NTR may participate in the formation of NFTs, another pathological character of Alzheimer's disease. When studying the brain of postmortem Alzheimer's disease patients, the ratio of p75NTR-positive neurons to the total number of neurons was found to be significantly higher in the CA1 and CA2 subfields of the hippocampus. Besides, a large proportion of these p75NTR expressing neurons are co-localized with Alz-50, whose antibody can be regarded as an indirect indication of hyperphosphorylated tau in the same subfields of Alzheimer's disease patients, suggesting that p75NTR may be involved in the formation of NFTs (Hu et al. 2002).

Our recent research demonstrated that deletion of the p75NTR gene can also exacerbate other Alzheimer's disease-type pathologies in the brain of Alzheimer's disease mice, such as microgliosis and microhemorrhage but does not reduce the memory deficits of Alzheimer's disease mice until 9 months of age (Wang et al. 2011).

## 5 Potential Use of p75NTR as a Therapeutic Target for Alzheimer's Disease

The roles of p75NTR in the pathogenesis of Alzheimer's disease described above strongly suggest that p75NTR represents a valid therapeutic target for the

treatment of Alzheimer's disease (Dechant and Barde 2002; Coulson et al. 2009; Knowles et al. 2009). However, as p75NTR has double-faced roles in A $\beta$  metabolism and neurotoxicity, it is cautioned that simply reducing expression of p75NTR may exacerbate Alzheimer's disease pathology, as knockout of the p75NTR gene results in increased A $\beta$  deposition and Alzheimer's pathologies in an APPswe/PS1dE9 transgenic mouse model. However, the use of the extracellular domain not only may attenuate neurotoxic signals by blocking A $\beta$ -p75NTR and proneurotrophin-p75NTR interactions on cell membranes but also may suppress A $\beta$  aggregation and deposition and enhance A $\beta$  clearance from the brain. In addition, it was found that the recombinant extracellular domain of p75NTR is also able to block the A $\beta$ - and proNGF-induced neurotoxicity in vitro (Wang et al. 2010a). This data suggests that recombinant extracellular domain of p75NTR may be a desirable anti-Alzheimer's disease agent with multiple functions to facilitate AB clearance by inhibiting AB aggregation, and to protect cholinergic neurons from A $\beta$ - and proneurotrophin-induced neurotoxicity by blocking their interaction with p75NTR, and to reduce A $\beta$  production by interrupt feedback loop between A $\beta$  and p75NTR (Fig. 1).

#### 6 Conclusion

p75NTR is an important molecule which has diverse functions linked with neuron development, aging, stress, and neuropathological conditions. Recent studies suggest that p75NTR plays a critical role in the pathogenesis, almost all hallmarks of Alzheimer's disease, starting from the generation of A $\beta$  and removal of A $\beta$  deposition, to mediate A $\beta$ -induced neuronal death, neurite degeneration, tau hyperphosphorylation, and cell cycle re-entry. The shedded p75NTR extracellular domain is regarded as an endogenous protective molecule to interrupt A $\beta$ /p75NTR feedback loop, block A $\beta$  neurotoxicity, and remove A $\beta$  deposits. Thus, the extracellular domain of p75NTR is a promising therapeutic target for the development of therapeutics for Alzheimer's disease.

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# Seizures Tip the Balance of Neurotrophin Signaling Toward Neuronal Death

Wilma J. Friedman, Audrey P. Le, Soyoung Choi, and Marta Volosin

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#### Abstract

Neurotrophins and proneurotrophins have opposing actions on neuronal survival, mediated by their respective receptors. Trk receptors are necessary for mature neurotrophins to support neuronal survival, while the p75 neurotrophin receptor ( $p75^{NTR}$ ) mediates proneurotrophin-induced apoptosis. Therefore, the consequences of neurotrophin actions depend on expression of the different neurotrophin receptor; regulation of proneurotrophin cleavage, which can determine which receptor is bound; and activation of downstream signaling events supporting neuronal survival or apoptosis. Several types of brain injury, including seizures, lead to altered levels of neurotrophin receptors, with increased neuronal expression of p75^NTR in the brain. Seizure-induced

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Audrey P. Le, present affiliation: Developmental and Regenerative Biology, Mt. Sinai School of Medicine, New York, NY, USA

Soyoung Choi, present affiliation: Department of Pathology and Laboratory Medicine, Weill Cornell Medical College, New York, NY, USA

W.J. Friedman (⊠) • A.P. Le • S. Choi • M. Volosin Department of Biological Sciences, Rutgers, The State University of New Jersey, Newark, NJ, USA

e-mail: wilmaf@andromeda.rutgers.edu; audrey.le@mssm.edu; soc2010@med.cornell.edu; marlapid@andromeda.rutgers.edu

injury also leads to increased secreted proNGF due to altered regulation of the extracellular proNGF-cleaving enzymes. Downstream signaling pathways triggered by proNGF activation of p75<sup>NTR</sup> include induction of the phosphatase PTEN, which antagonizes TrkB signaling triggered by BDNF and prevents the activation of Akt. Thus, the increase in extracellular proNGF and elevated levels of p75<sup>NTR</sup> that occur after injury create a milieu that leads to neuronal death.

#### **Keywords**

Brain-derived neurotrophic factor • Hippocampus • Nerve growth factor • p75 Neurotrophin receptor • Trk

## 1 Introduction

Neurotrophins have been well established as trophic factors in the peripheral and central nervous systems, supporting neuronal survival and promoting differentiation (Barde 1994; Huang and Reichardt 2001). The neurotrophin family, consisting of NGF, BDNF, NT3, and NT4, binds to two distinct types of receptors: a member of the Trk receptor tyrosine kinase family and the p75 neurotrophin receptor (p75<sup>NTR</sup>). The role of Trk receptors in mediating neuronal survival, differentiation, and synaptic function has been well defined, and the major signaling pathways activated include the PI<sub>3</sub> kinase-Akt, ras-MAP kinase, and PLC $\gamma$  signaling pathways, which have been extensively studied (Friedman and Greene 1999; Kaplan and Miller 2000; Patapoutian and Reichardt 2001). In contrast, the p75<sup>NTR</sup> has been more of an enigma, since it has been implicated in mediating many diverse cellular functions such as survival (Hamanoue et al. 1999; Chu et al. 2007), apoptosis (Casaccia-Bonnefil et al. 1996; Frade et al. 1996; Coulson et al. 1999), and axonal growth (Yamashita et al. 1999; Sun et al. 2012), depending upon the cellular context. Following numerous different types of injury, including seizures, traumatic brain injury, and axotomy, p75<sup>NTR</sup> has been shown to signal apoptosis (Friedman 2000; Beattie et al. 2002; Troy et al. 2002; Harrington et al. 2004). Although p75<sup>NTR</sup> can bind all the neurotrophins with low affinity, uncleaved proneurotrophins bind  $p75^{NTR}$  with high affinity (Lee et al. 2001) due to the association with the coreceptor sortilin (Nykjaer et al. 2004) and potently induce neuronal death (Harrington et al. 2004; Volosin et al. 2008). Thus, neurotrophins can signal survival or apoptosis, depending on which form of the factor is secreted, which can determine which receptor and signaling pathways are activated. Therefore, determination of neuronal survival or death following injury depends on several events, such as altered expression of the different neurotrophin receptors, p75<sup>NTR</sup> and the Trk family. Moreover, regulation of proneurotrophin processing to determine whether the cleaved or uncleaved form is secreted and stabilized in the extracellular milieu can be a critical determinant of which receptor is engaged, leading to neuronal survival or death. Thus, a delicate balance of receptor expression and downstream signaling, as well as proneurotrophin processing, can critically determine the outcome for signaling neuronal survival or death.

# 2 p75<sup>NTR</sup> and Apoptotic Signaling

Many injury conditions, including seizures, lesions, and traumatic injuries, have been shown to induce the  $p75^{NTR}$ , suggesting an important role in mediating neuronal responses to injury (Ernfors et al. 1989; Roux et al. 1999). Moreover, in mice lacking this receptor, neuronal death is significantly reduced following injury (Beattie et al. 2002; Troy et al. 2002). Although  $p75^{NTR}$  is a member of the TNF receptor family of death receptors, it does not signal through the same pathways as other members of this family, which generally recruit adapter proteins to the trimerized receptor to activate caspase-8 and the extrinsic apoptotic pathway (Green 2003). In contrast, activation of  $p75^{NTR}$ -mediated apoptosis requires phosphorylation of c-Jun N-terminal kinase (JNK) (Yoon et al. 1998; Friedman 2000) and phosphorylation of the pro-apoptotic proteins Bad (Bhakar et al. 2003) and BimEL (Becker et al. 2004), eliciting the release of cytochrome c from the mitochondria and the subsequent activation of caspase-9, caspase-6, and caspase-3 (Wang et al. 2001; Troy et al. 2002). Thus, apoptotic signaling via  $p75^{NTR}$  is different from the classic members of the TNFR family of death receptors.

The p75<sup>NTR</sup> has no kinase activity and signals by recruiting intracellular binding proteins to activate downstream events. The receptor undergoes regulated intramembrane proteolysis (RIP), generating a C-terminal fragment (CTF) and ultimately the intracellular domain (ICD) that interacts with intracellular proteins such as NRIF, NRAGE, and NADE to promote cell death (Barker 2004), ultimately activating the intrinsic caspase cascade. Genetic mutation analysis of different domains of the p75<sup>NTR</sup> has demonstrated that specific amino acids distributed throughout the death domain are required for the activation of the caspase pathway, leading to cell death, while other domains are necessary for activation of RIP2 binding, leading to NF-KB activation and cell survival (Charalampopoulos et al. 2012). Activation of NF- $\kappa$ B by p75<sup>NTR</sup> has been observed in Schwann cells (Carter et al. 1996), but not in hippocampal neurons or astrocytes (Volosin et al. 2008). The activation of distinct cell-specific signaling pathways is likely regulated by the expression or recruitment of the specific adapter proteins that link the receptor to the different downstream pathways. The scattered amino acid residues that are required for caspase-3 activation suggest that multiple binding proteins are necessary for activation of this pathway, possibly forming a multi-molecular complex binding to the receptor.

Following damage in the brain, numerous cytokines and trophic factors are released that can impact upon the injured neurons. Mature neurotrophins, such as BDNF, are present in the hippocampus and can activate its TrkB receptor to promote signaling via the PI3K-Akt and ras-ERK pathways to support survival of these neurons. The ability of neurotrophins to support neuronal survival can be affected by changes in the neurotrophin receptor levels and signaling that occur following seizures. Seizures, as well as other types of injury, lead to an induction of p75<sup>NTR</sup> expression in the hippocampus (Roux et al. 1999; Troy et al. 2002). A previous study has shown that along with the increase in p75<sup>NTR</sup>, there is a concomitant decrease in TrkB (Unsain et al. 2008), tilting the balance of signaling towards neuronal death. The induced p75<sup>NTR</sup> on hippocampal neurons can be activated by proNGF, which is

released following injury, leading to neuronal death. Neurons that may express both p75<sup>NTR</sup> and TrkB and may be exposed to BDNF and proNGF must integrate these signals to a decisive outcome, survival, or death. Interestingly, in contrast to sympathetic neurons where activation of Trk prevents p75<sup>NTR</sup>-mediated apoptosis (Miller and Kaplan 2001), CNS neurons expressing both p75<sup>NTR</sup> and TrkB, and exposed to both proNGF and BDNF, were not protected from cell death by the BDNF. This was due to the induction of the PTEN phosphatase by p75<sup>NTR</sup> activation, which prevented the ability of TrkB to activate Akt, thereby inhibiting the ability of BDNF to promote survival and facilitating the apoptotic signaling of p75<sup>NTR</sup> (Song et al. 2010). When the induction of PTEN by proNGF was prevented, BDNF was able to rescue hippocampal neurons from death following seizures, indicating that the induction of PTEN is a critical checkpoint in the balance between survival and apoptotic signaling (Song et al. 2010). Thus, induction of neuronal apoptosis by proNGF in the presence of mature neurotrophins requires the suppression of Trk signaling as well as the activation of JNK and subsequent caspase cascade.

# 3 Seizure-Induced Changes in Connectivity

A major consequence of epileptic seizures is the dendritic sprouting and synaptic reorganization that leads to the formation of aberrant connections and the development of spontaneous recurrent seizures (Williams et al. 2009). Many studies have shown that BDNF is elevated in the hippocampus following different types of seizures (Ballarin et al. 1991). Levels of the TrkB receptor are particularly high in the mossy fibers, projections that undergo sprouting and circuit reorganization, which are thought to participate in the development of recurrent seizures, leading to increased seizure susceptibility (Koyama et al. 2004; McNamara and Scharfman 2012). In contrast to the stimulation of axonal and dendritic sprouting induced by BDNF, recent studies have demonstrated that p75<sup>NTR</sup> can induce axonal degeneration. In fact, the activation of caspase-6 by p75<sup>NTR</sup> has been implicated not only in cell death but also in axonal degeneration (Park et al. 2010), suggesting that activation of p75<sup>NTR</sup> may play a broader role following injury, not only inducing neuronal loss but also potentially triggering more subtle but widespread responses involving axonal loss and circuit remodeling. Thus, in addition to having opposing actions with respect to neuronal survival, TrkB and p75<sup>NTR</sup> have opposing actions in regulating axonal growth, with TrkB eliciting sprouting and  $p75^{NTR}$  evoking degeneration, both of which may come into play following seizures with the consequent formation of aberrant circuits.

# 4 Regulation of p75<sup>NTR</sup> Following Injury

The p75<sup>NTR</sup> is not widely expressed in the adult brain but is induced following numerous different types of insults, including seizures, axotomy, and trauma. This receptor is also upregulated in neurodegenerative diseases such as Alzheimer's

disease. The induction of this receptor in all these pathological situations raises the question of whether there are common signals present in all these conditions that regulate expression of  $p75^{NTR}$ . One common feature of all these injury and disease conditions is hypo-osmolarity, which has been shown to regulate  $p75^{NTR}$  expression via activation of Sp1 transcription factors (Ramos et al. 2007). Another common characteristic of these conditions is the presence of inflammation in the brain and high levels of inflammatory cytokines such as IL-1 $\beta$  and TNF $\alpha$ . These cytokines can induce expression of  $p75^{NTR}$  in hippocampal neurons in vitro (Choi and Friedman 2009) and in vivo and may contribute to the mechanism by which  $p75^{NTR}$  is induced on vulnerable neurons to promote neuron loss. IL-1 $\beta$  is known to enhance neuronal vulnerability to many types of damage, including traumatic brain injury and stroke. In culture, preexposure of hippocampal neurons to IL-1 $\beta$  enhances the apoptotic response to proNGF, suggesting that proNGF is part of the mechanisms by which IL-1 $\beta$  increases neuronal vulnerability.

#### 5 Regulation of Extracellular Proneurotrophin Stability

Neurotrophins are synthesized as precursor proteins that undergo cleavage in the trans-Golgi network by enzymes such as furin and other proconvertases with the cleaved product being secreted as the mature neurotrophin (Seidah et al. 1996). As the precursors of neurotrophins, proneurotrophins are transiently present in neurons and glial cells in the brain; therefore, detection of intracellular proneurotrophins does not indicate which form of the proteins is secreted. The mechanisms by which proneurotrophins may be secreted without being processed are undefined; however, since neurotrophins and proneurotrophins can have opposing actions in mediating neuronal survival or death, and axonal sprouting and degeneration, it is clearly critical to maintain tight regulation of proneurotrophin cleavage. Therefore, in addition to the intracellular processing of this protein by furin, mechanisms also exist for promoting proneurotrophin processing in the extracellular environment. Extracellular enzymes that can cleave proneurotrophins include matrix metalloproteinases (MMPs) and the tissue plasminogen activator (tPA)-plasminogen cascade (Lee et al. 2001). In the pentylenetetrazole (PTZ) model, seizures increased levels of MMP-9, promoting the conversion of proBDNF to BDNF (Mizoguchi et al. 2011). In addition, electroconvulsive seizures (ECS) increased levels of proconvertase 1 and tPA (tissue plasminogen activator), which also promote processing of proBDNF to BDNF (Segawa et al. 2012). Thus, there is evidence that seizures promote cleavage of proBDNF to BDNF, which then increases the likelihood of recurrent seizures (McNamara and Scharfman 2012). However, in contrast to proBDNF, the unprocessed proNGF has been detected in cerebrospinal fluid under conditions of injury and neurodegenerative disease (Harrington et al. 2004; Le and Friedman 2012), indicating that the NGF precursor protein was secreted without being cleaved and is stable in the extracellular environment and not cleaved by extracellular proteases. This indicates that alterations in the levels and/or activity of these cleaving enzymes occur as a consequence of the injury leading to stabilization of the proneurotrophin. Unlike proBDNF, proNGF is not cleaved by MMP9 but can undergo extracellular cleavage by MMP7. Seizure-induced injury leads to decreased levels of MMP7 protein and activity and increases in tissue inhibitor of metalloproteinase (TIMP)-1 (Rivera et al. 1997), resulting in inhibition of proNGF cleavage (Le and Friedman 2012). Moreover, infusion of exogenous MMP7 into the hippocampus following seizures resulted in cleavage of extracellular proNGF and provided neuroprotection (Le and Friedman 2012). Thus, the altered balance of these extracellular enzymes and their endogenous inhibitors regulates the stability of proNGF in the extracellular milieu, providing a specific ligand for binding the injury-induced p75<sup>NTR</sup> and promoting neuronal loss following seizures.

## 6 Conclusion

A tightly regulated balance exists between the ability of neurotrophins to promote neuronal survival and apoptosis. In the healthy brain, proneurotrophins are

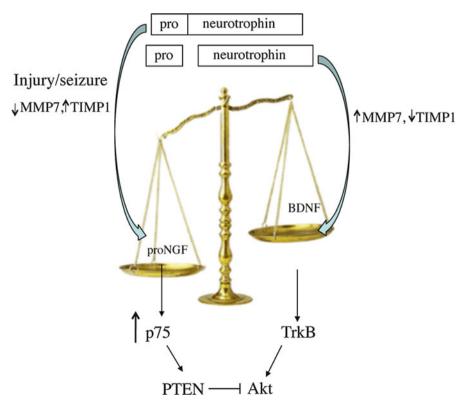


Fig. 1 Tilting the balance: seizure-induced alterations in the balance of proneurotrophin/ neurotrophin signaling that lead to neuronal death

processed to generate the mature trophic factors, which signal via their Trk receptors to influence neuronal, survival differentiation and function. Following seizures and other types of injury, the induction of p75<sup>NTR</sup> expression on injured neurons and extracellular presence of proNGF shift the balance of neurotrophin actions towards inducing apoptosis of the injured cells (Fig. 1).

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# Synthesis, Trafficking and Release of BDNF

Jian-Jun Lu, Miao Yang, Ying Sun, and Xin-Fu Zhou

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#### Abstract

Brain-derived neurotrophic factor (BDNF), a member of the neurotrophin family, plays an important role in neuronal survival, differentiation, neurite outgrowth, spine formation, synaptic plasticity, and memory. The regulation of its biological activity is well controlled by its gene expression, axonal transport, and release. It is well known that BDNF is synthesized and released in an activity-dependent manner. However, the molecular mechanism for its processing, axonal transport, and release is still not completely understood. In our recent studies we found that Huntingtin-associated protein (HAP1) and sortilin play important roles in BDNF trafficking and processing. HAP1 plays critical roles in BDNF intracellular trafficking, dendritic targeting, and metabolism by binding to the prodomain of

J.-J. Lu (🖂)

M. Yang • Y. Sun • X.-F. Zhou

e-mail: miao.yang@unisa.edu.au; Annie.Sun@unisa.edu.au; xin-fu.zhou@unisa.edu.au

Division of Health Sciences, University of South Australia, Adelaide, SA, Australia e-mail: steven.lu@unisa.edu.au.

School of Pharmacy and Medical Sciences, Faculty of Health Sciences, University of South Australia, Adelaide, SA, Australia

BDNF and forming a complex with sortilin. In this chapter, we focus on the recent progress made in understanding the molecular mechanism underlying the biosynthesis, transport, and release of BDNF, emphasizing the role of HAP1 and sortilin in the intracellular trafficking of BDNF.

List of Abbreviations				
ARF	ADP-ribosylation factor			
BDNF	Brain-derived neurotrophic factor			
BDNF-GFP	Green fluorescent protein-tagged BDNF			
CaRF	Calcium-responsive transcription factor			
CREB	cAMP-responsive element-binding protein			
HAP1	Huntingtin-associated protein 1			
HFS	High-frequency stimulation			
LFS	Low-frequency stimulation			
LTD	Long-term depression			
LTP	Long-term potentiation			
MMPs	Matrix-metalloproteinases			
MPR	Mannose 6-phosphate receptors			
proBDNF	Precursor of brain-derived neurotrophic factor			
RGCs	Retinal ganglion cells			
SNP	Single nucleotide polymorphism			
TGN	Trans-Golgi network			
tPA	The tissue-plasminogen activator plasmin			
trkB	Tyrosine kinase receptor B			

## 1 Introduction

Brain-derived neurotrophic factor (BDNF), a member of the neurotrophin family, plays important roles in neuronal differentiation and survival, synaptic plasticity, and long-term potentiation during the entire life. BDNF mainly expresses in neural tissues and it is transported both anterogradely and retrogradely in neurons. It has an autocrine and paracrine effect on neuronal cells. BDNF mediates its biological action through binding to its high affinity receptor, tyrosine kinase receptor B (trkB), and p75, the pan receptor for other members of the neurotrophin family. However, the precursor of BDNF (proBDNF) has different binding characteristics and distinct biological activity in comparison with mature BDNF. It is well known that the biological action of BDNF is activity dependent and has been extensively explored. However, the molecular mechanism of its axonal transport and processing is not completely understood. Here in this chapter, we focus on BDNF axonal transport and describe the recent progress in BDNF trafficking, processing, and release.

#### 2 The Expression of BDNF

The BDNF expression starts at an early embryonic stage. Its mRNA is expressed throughout development reaching the highest levels by postnatal day. In adult animals and humans, BDNF is expressed in the brain, with the highest levels in the hippocampus and cortex, suggesting an important role of this factor in the maintenance of the normal central nervous system. In humans, the BDNF gene is located in chromosome 11.

Like all neurotrophins, BDNF is expressed as a precursor encompassing two domains, the prodomain and the mature domain. The BDNF coding sequence has been determined in mice, pigs, rats, and humans. In all of these species, the mature BDNF shows complete identity in amino acid sequence. The expression of BDNF is well controlled in neurons by transcriptional regulation and differential targeting of BDNF transcripts at distinct cellular locations. In addition, neurotransmitters and membrane depolarization are also involved in the regulation of BDNF expression.

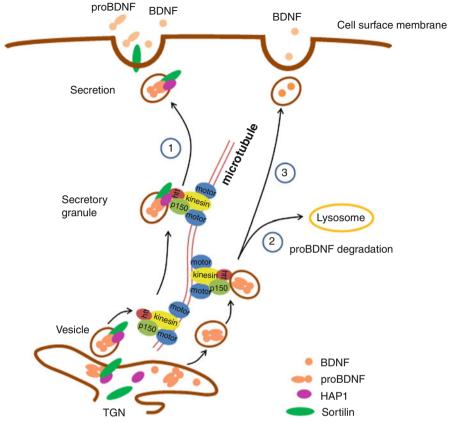
BDNF has at least 18 transcripts produced by alternative promoters, splicing, and polyadenylation sites, but each encodes an identical BDNF protein product. This complex transcriptional organization seems to provide multiple layers of regulation through alternative promoter usage, differential mRNA stability, and differential subcellular localization of either mRNA or protein. For example, it has been shown that the expression of the BDNF gene is regulated by neural activity through calcium-mediated pathways and that different BDNF mRNA transcripts driven by different promoters are differentially regulated. BDNF exon I and exon IV transcripts have previously been characterized as the most highly induced BDNF mRNAs, in response to excitatory treatment and KCI-mediated membrane depolarization in embryonic cortical neuron cultures. Several calcium-responsive elements and transcription factors binding to these elements in the promoter regions have been characterized. BDNF exon V, exon VII, exon VIII, and exon IXA transcripts have also been shown to be regulated by kainic acid and that the induction magnitude is comparable to that of BDNF exon I and IV transcripts. Importantly, differential regulation of BDNF mRNAs is found in mood disorder, exercise, and learning. Thus, it is suggested that the differential regulation of BDNF exon mRNAs might be involved in different neurodegenerative diseases in which BDNF levels are altered.

Regulation of the BDNF gene has been well studied. In addition to the transcriptional regulation, different BDNF messenger RNA (mRNA) transcripts are found at different subcellular locations in hippocampal and cortical neurons. BDNF expression was remarkably increased in the neocortex following experimentally induced seizures (Ernfors et al. 1991; Isackson et al. 1991). Several other studies also reported regulation of BDNF in vivo by sensory stimulation and in vitro by neuronal activity (Castren et al. 1992; Bozzi et al. 1995; Ghosh and Greenberg 1995; Lauterborn et al. 1996), suggesting that this is an important mechanism by which levels of BDNF are normally regulated. The transcription of BDNF can be regulated by neuron activity in hippocampal neurons (Tongiorgi et al. 1997). A recent study also showed that strong depolarizing stimuli, both in vitro and in vivo, cause accumulation of BDNF mRNA and protein in the distal dendrites, through the activation of the *N*-methyl-D-aspartate and tyrosine kinase receptor B (TrkB) (Tongiorgi et al. 2006). Indeed, neuron activity can modulate the expression of many genes that affect the synaptic plasticity (Prakash et al. 2008). Therefore, knowledge regarding gene regulation of BDNF is important, in order for us to understand how altered BDNF expression and release can have an impact on the function of the neuronal circuit. Future characterization of the regulatory sequences and transcription factors mediating regulation of novel BDNF transcripts in different disease models is important for understanding BDNF gene regulation and its contribution to pathology.

### **3 BDNF Processing**

BDNF, like other neuropeptide, is initially synthesized as a precursor containing a signal peptide in the endoplasmic reticulum. The signal peptide is cleaved off immediately, yielding so-called proproteins which undergo further posttranslational modifications in Golgi apparatus. Following the cleavage of the signal peptide, proBDNF is packaged in trans-Golgi network (TGN) for sorting into secretory vesicles. However, proBDNF may be converted into mature BDNF via both intracellular and extracellular enzymes. Studies show that the cleavage of proBDNF on the RVRR motif occurs in the trans-Golgi, by members of the subtilisin-kexin family of endoproteases such as furin, or in the immature secretory granules by proprotein convertases. It is also observed that tissue-plasminogen activator (tPA) converts proBDNF to the mature BDNF by activating the extracellular protease plasmin, which is critical for long-term potentiation (LTP) of the mouse hippocampus. Whether the conversion of proBDNF to mature BDNF is regulated by extracellular signals is not yet known. To understand the mechanisms that regulate the conversion of proBDNF to mBDNF, some experiments were undertaken to investigate how the physiologically relevant condition is associated with the secretion of specific isoforms of endogenous BDNF. It was demonstrated that under conditions that induce long-term depression (LTD), proBDNF is mainly secreted.

A stable status of endogenous proBDNF is important for its efficient axonal transport. It has been identified that sortilin binds to the prodomain of BDNF, which modulates its stabilization and regulates it into the secretory pathway. This regulation is dependent on the function of sortilin domains. Deletion of sortilin intracellular domain causes BDNF missorting to the constitutive secretory pathway, while overexpression of sortilin ectodomain impairs lysosomal targeting and leads to degradation of BDNF. Our recent data shows that co-expression of transfected proBDNF, HAP1, and sortilin in HEK293 cells can prevent proBDNF degradation, whereas the absence of sortilin significantly increased proBDNF degradation. The effect of sortilin on stabilization of proBDNF protein is dose dependent. These results suggest that normal BDNF trafficking and stabilization in neurons is dependent on the intact function of sortilin, in particular the complex of proBDNF/HAP1 and sortilin (Fig. 1).



proBDNF anterograde transport

**Fig. 1** proBDNF anterograde transport. New synthesized proBDNF proteins exist from endoplasmic reticulum (ER) and move to trans-Golgi network (TGN) for packaging and sorting. During the process, proBDNF can form a complex with HAP1 and sortilin and interacts with kinesin motor proteins through Htt and p150Glued. This proBDNF/kinesin complex is transported along with the microtubule to axonal terminal for secretion (I). When proBDNF is packaged itself in a vesicle without the participation of HAP1 and sortilin and is carried by the kinesin complex, some proBDNF proteins can be sent to lysosome for degradation (2), or a part of proBDNF can be converted to BDNF, which is sent to axonal terminal for secretion (3)

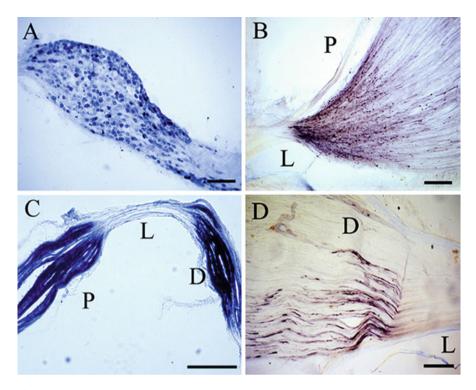
#### 4 Transport of BDNF

The classic neurotrophic hypothesis is that neurotrophins are synthesized in peripheral target tissues, uptaken by the innervating nerve terminals, and retrogradely transported to the neuronal cell body, subsequently exerting their biological function by binding and activating their specific receptors. Like other neurotrophic factors, BDNF was initially considered to be a retrogradely transported neurotrophic factor. <sup>125</sup>I-labelled BDNF injected into peripheral target tissues was found retrogradely transported into the subpopulation of peripheral and central neuronal cells. A study by intrastriatal infusion of BDNF revealed its retrograde transport in neurons in the brain. Furthermore, intraparenchymal injections of BDNF resulted in widespread retrograde transport throughout the central nervous system (CNS). However, the retrograde mode of BDNF transport cannot explain some of its functional roles. Moreover, it has been demonstrated that BDNF is produced in the sensory neurons because its mRNA and protein are both expressed in sensory neurons. Conversely, BDNF mRNA is not expressed in peripheral axonal terminals, whereas BDNF protein is detectable in terminals, suggesting that BDNF might be anterogradely transported in neurons.

We are among the first few groups that discovered the anterograde transport of BDNF in sensory neurons. We demonstrated that BDNF is transported anterogradely via both peripheral and central processes of spinal sensory neurons. Using specific antisera, we examined the distribution of BDNF immunoreactivity and found it to be present within a subpopulation of sensory neurons, primarily those with a small-to-medium diameter. The immunoreactivity was accumulated on both the distal and proximal sides of a ligature on the sciatic nerve (Fig. 2). The accumulation on the distal side, but not on the proximal side, was substantially reduced by pretreatment with BDNF antibodies in vivo. In contrast to the periphery, the immunoreactivity was only accumulated on the proximal side of a lesion of the dorsal root. In the spinal cord, most nerve terminals immunoreactive for BDNF were identified in lamina II. Lesion of the dorsal root led to a reduction of these nerve terminals. These studies have provided clear evidence that the factor is transported not only retrogradely, but also anterogradely from the spinal ganglia to terminals in the periphery and spinal cord. Consistently, studies from other research groups also confirmed anterograde transport of BDNF.

It was shown that BDNF is widely distributed in nerve terminals, even in brain areas such as the striatum that lack BDNF messenger RNA, suggesting this factor is anterogradely supplied to some areas of the brain. Interestingly, by using the same method, a study in our laboratory also showed that proBDNF is both anterogradely and retrogradely transported in sensory neurons. Similar to the mature BDNF, after ligation of the sciatic nerve, increased proBDNF expression can be detected both proximal and distal to the site of ligation. In CNS, analysis of distribution of BDNF and mRNA revealed more detailed transport patterns of BDNF in some neural structures. BDNF is also anterogradely transported in hippocampus mossy fibers and nerve terminals in other brain structures. Furthermore, BDNF is also found to be anterogradely transported in retina cells. Retinal ganglion cells (RGCs) also anterogradely transport exogenous neurotrophins including BDNF to the midbrain tectum/superior colliculus with significant downstream effects.

Consistent with in vivo studies, in vitro studies also showed BDNF is transported in an anterograde mode. Several neuronal types can internalize exogenous neurotrophic factors which then can be anterogradely transported to axonal



**Fig. 2** Microphotographs showing the immunoactivity and transport of BDNF in the peripheral nervous system of adult rats. Panel (a): nodose ganglion; panel (b): proximal segments of ligated sciatic nerve; panel (c): ligated vagal nerve; (d): distal segment of ligated sciatic nerve. Abbreviations: *L* ligation site, *P* proximal to the ligation site, *D* distal to the ligation site. Scale bars in (a), (b), and (d) = 100  $\mu$ m, in (c) = 1 mm

pathways. Neuronal bodies and dendrites possess receptors for the ligands, and a significant amount of internalized ligands are targeted for anterograde axonal transport, apparently with subsequent axonal release. While numerous studies demonstrate anterograde transport of trophic factors in axons, only few provide direct evidence of transcytosis of endogenous or exogenous trophic factors. Interestingly, a study comparing the trafficking pattern of BDNF in axons and dendrites demonstrated that axonal anterograde transport is the major form of BDNF transport. However, the axonal transport pattern is different from the dendritic transport pattern. The pattern and velocity of the trafficking of fluorescent protein-tagged BDNF is different between axons and dendrites. The movement of BDNF-GFP puncta in axons is faster than in dendrites, suggesting that the anterograde transport in axons may be the dominant site of BDNF release.

The anterograde transport of BDNF appears to be a universal phenomenon in the mammal nervous system. However, the real-time in vivo record of endogenous BDNF transport is lacking because of technique difficulties and a relatively small amount of detectable endogenous BDNF. The anterograde mode of BDNF

transport is critical for the neurotropism, which is not merely an alternative way for retrograde transport. Numerous studies have demonstrated that the bidirectional transfer and trafficking of BDNF through the neural network is a fundamental mechanism for its trophic regulation of neuronal survival, differentiation of phenotypes and dendritic morphology, synaptic plasticity, as well as excitatory neurotransmission. It is now clear that newly synthesized BDNF is packaged into vesicles and anterogradely transported to neuronal terminals for secretion, via interaction with kinesin motor protein complex along the microtubules. The molecular mechanism underlying BDNF intracellular trafficking still remains to be elucidated. Nevertheless, recent studies have provided some evidence of HAP1 and sortilin playing critical roles in BDNF trafficking and processing.

In search of the potential molecules involved in BDNF intracellular trafficking, we found that HAP1 and sortilin are ideal candidates. HAP1 is a cytoplasmic protein enriched in the brain and spinal cord. HAP1 was identified as a Huntingtin (Htt)-binding partner by yeast two-hybrid screens. It also associates with microtubules in addition to several vesicular organelles. Htt is a scaffold protein predominantly found in the cytoplasm, where it associates with various vesicular structures and molecular motors to form a cargo complex and play a role in intracellular trafficking. Despite the important role of HAP1 in neurons, the precise function of HAP1 in neurons remains to be seen. Some recent studies have shown that HAP1 is involved in endocytosis and vesicular trafficking. The expansion of polyglutamine (polyO) repeats in Htt and the deletion of either HAP1 or p150Glued can reduce BDNF axonal transport and lead to the degeneration of striatal neurons. It is suggested that a molecular bridging of HAP1 with BDNF and kinesin motor complex may be involved in BDNF axonal transport. For example, HAP1 interacts with the p150Glued subunit of dynactin and regulates axonal trafficking. HAP1 also mediates the anterograde transport of the GABAA receptor to synapses and of neurotrophic factor receptors for neurite outgrowth.

Sortilin is a member of the Vps10p domain receptor family including SorLA and SorCS1, SorCS2, and SorCS3-3. Sortilin is highly expressed in the cortex and hippocampus. It can function as a cell surface receptor for proneurotrophins, neurotensin, and lipoprotein lipase. Sortilin is mainly distributed in intracellular compartments where it binds to diverse ligands including hydrolases. Sortilin is a sorting molecule which sorts lysosomal hydrolytic enzymes to lysosomes and BDNF into secretory pathways. Its short (53 aa) cytosolic tail contains a motif that is similar to that of the sorting receptor mannose 6-phosphate receptors (MPR) and low-density lipoprotein (LDL) receptors, which interact with the family of Golgi-localizing,  $\gamma$ -adaptin ear homologous ADP-ribosylation factor (ARF)binding proteins (GGA) that shuttle between the Golgi and endosomes/lysosomes. The C-terminal acidic cluster-dileucine motif mediates endocytosis of the receptor. The cytosolic domain of sortilin also interacts with the retromer complex, which recycles sortilin back to Golgi for further trafficking and sorting of proteins. The retromer complex is composed of 3–4 VPS molecules VPS35, VPS29, and VPS26.

There are other factors which may directly or indirectly affect the BDNF trafficking. A Val66Met substitution of BDNF led to a significant decrease

of BDNF trafficking and secretion and was found to be associated with increased susceptibility to neuro-disorders including Alzheimer's disease, Parkinson's disease, depression, and bipolar disorder. Since sortilin interacts with the prodomain of BDNF in a discrete region containing the Met substitution, the altered BDNF trafficking and secretion is likely due to the disruption of the interaction between BDNF and sortilin. This concept is also supported by the evidence that sortilin is involved in anterograde transport of BDNF. These studies reveal a role of sortilin in mediating trkB receptors and BDNF anterograde transport, which is important for neurotrophin-induced neuronal survival. In addition, Jun NH (2)-terminal kinase-interacting protein 3 (JIP3) mediates trkB axonal anterograde transport and enhances BDNF signaling by directly interacting with trkB juxtamembrane region and linking trkB to kinesin-1. Given that both JIP3 and sortilin bind to trkB, their effect on enhancing BDNF signaling may rely on functions of both proteins. Nonetheless, these findings provide a novel mechanism of how the TrkB anterograde transport can be coupled to BDNF signaling in axons.

Our recent studies demonstrate that HAP1 and sortilin may play an important role in BDNF trafficking and processing. We investigated the HAP1 and BDNF interaction and the role of HAP1 in BDNF trafficking by using cell transfection and HAP1 knockout mice. Our mapping studies demonstrate that the region of HAP1 between amino acids 371 and 445 contains proBDNF-binding sites. Pull-down assays show that the region between amino acids 65 and 90 of the BDNF prodomain has a key structure for HAP1 binding. Our results also show that HAP1 regulates axonal transport and the activity-dependent release of BDNF, through HAP1 interaction with proBDNF but not mature BDNF. Reduced association of HAP1 with proBDNF by polyQ-expanded Huntingtin or Val66Met BDNF variant can negatively impact on proBDNF trafficking. We also observed a similar reduction in HAP1 and proBDNF interaction using the brain homogenate of Huntington's disease patients, suggesting that this reduced association of HAP1 with proBDNF can occur in vivo and may cause a deficiency of BDNF anterograde axonal transport. This hypothesis is supported by the evidence that ligation of the sciatic nerve did not show increased accumulation of BDNF in the proximal and distal segments of the ligation site and the deficit in proBDNF trafficking in HAP1-/- neurons was rescued by introduction of HAP1 cDNA in these neurons. Therefore, our data suggests that HAP1 may play a regulatory role in axonal transport of BDNF. Recently we also found that HAP1 can interact with proBDNF and form a complex with sortilin. This complex is found in most cellular organelles involved in protein sorting and trafficking such as ER, TGN, and Golgi complex. The complex can stabilize the proBDNF. In co-transfected HEK 293 cells, the complex helps to prevent proBDNF degradation. Furthermore, the complex facilitates furin cleavage to release mature BDNF.

#### 5 BDNF Release and Secretion

There are two separate types of secretion pathways for BDNF: the regulated pathway and the constitutive pathway. The former employs  $Ca^{2+}$ -dependent exocytosis of

secretory granules, while the latter employs a distinct type of vesicles which releases the cargo by default when reaching the plasma membrane. It is suggested that the subcellular location of BDNF is important for understanding the function of BDNF and its role in synaptic plasticity. Since the level of endogenous BDNF is very low in most neuronal tissues, overexpression of BDNF-GFP has been used to study the transport and release of BDNF in neuronal culture. Although the expression of tagged protein may have altered binding to sorting partners, the distribution of BDNF-GFP is similar to that observed for endogenous BDNF. In dendrites, BDNF-GFP is packaged into secretory granules for the regulated secretion pathway, suggesting a dendritic secretion of BDNF. This finding is further supported by the evidence that local synthesis of dendritic BDNF and dendritic BDNF mRNA transcript has been detected in cultured hippocampal neurons. Also, alteration of dendritic BDNF mRNA can cause the change of the local BDNF level. However, a recent study, using specific monoclonal antibodies to label proBDNF and mature BDNF, has shown that proBDNF and mature BDNF exist in presynaptic dense core vesicles but not in dendrites. Hippocampal neurons from mutant mice with overexpressed BDNF also show that BDNF is only present in presynaptic dense core vesicles. The finding challenges the previous concept of activity-dependent dendritic synthesis and release of BDNF. However, the finding supports the conclusion that BDNF is transported in an anterograde mode, which is in contrast to the previous concept that BDNF is retrogradely transported in target cells. This in vivo study appears to provide convincing evidence that BDNF is located in presynaptic dense core vesicles.

There are controversial views about the neuronal release of proBDNF. Barde's group reported that proBDNF is only an intermediate metabolic product of precursor BDNF, which only presents transiently and does not have biological functions. In contrast, a similar study claims that proBDNF is present and is released from neurons exerting biological functions. In this study, different culture conditions were applied to avoid possible glial contamination, and alpha 2 antiplasmin was used to reduce the cleavage of secreted proBDNF. The secretion of proBDNF from cultured hippocampus neurons was convincingly demonstrated. The conclusion drawn from this study appears to be consistent with that from a previous study, reporting that the cleavage of proBDNF by tPA/plasma is essential for long-term hippocampus plasticity. tPA converts proBDNF to mature BDNF by activating the extracellular protease plasmin, and such conversion is critical for L-LTP expression in mouse hippocampus. Moreover, application of mBDNF is sufficient to rescue L-LTP when protein synthesis is inhibited, which suggests that mBDNF is a key protein synthesis product for L-LTP expression.

To reveal which isoform of BDNF is secreted under physiological stimulation, frequency stimulation of neuronal cells was performed. Studies have shown that proBDNF is mainly secreted from neuronal terminals during low-frequency stimulation (LFS). In contrast, mature BDNF is the major secreted isoform when neurons were stimulated with high-frequency stimulation (HFS, a condition that induces LTP). Thus, the secretion of either proBDNF or mature BDNF is dependent on low- or high-frequency neuronal activity. These findings suggest that high-frequency neuronal activity can enhance the function of mature BDNF.

In addition, the secretion of proBDNF or mature BDNF is also controlled by activitydependent synaptic competition/elimination. ProBDNF can trigger synaptic depression and subsequent retraction of the axonal terminal through p75 receptors in Xenopus neuromuscular synapses. Active axonal terminals can cleave proBDNF to mature BDNF which protects the active terminal from depression and retraction.

BDNF release is a regulated response to its functional requirement. The dendritic BDNF synthesis and the release of BDNF-GFP under high-frequency electric stimulation provide evidence that BDNF release is in an activity-dependent manner from dendrites and axons. Although the importance of BDNF in synaptic plasticity has been clearly established, direct evidence for the activity-dependent dendritic secretion of BDNF has not been obtained. In the cerebellum, the depolarizationdependent release of BDNF from granule cells was reported to be regulated by the CAPS2 molecule, which leads to cell differentiation and survival during cerebellar development. Furthermore, exercise can increase the serum level of BDNF in human subjects. The molecular interaction has been determined to be responsible for BDNF-regulated release.

## 6 BDNF Transport and Release in Neurological Diseases and Neurotoxicity

BDNF is associated with the development of many neurological and psychiatric disease conditions because of its wide distribution and great functional diversities. Deficiencies in its synthesis, trafficking, and release play important roles in the pathogenesis of a wide range of neurodegenerative diseases and cognitive disorders. However, the molecular dissection of the trafficking and release of BDNF in diseased condition is limited, although the axonal transport in these disease conditions has been extensively studied. Axonal transport defects have been described as an early pathological feature in a variety of animal models of Alzheimer's disease (AD) and tauopathies. It is clear that BDNF axonal transport of BDNF is impaired in AD. In Huntington disease, there is reduced expression of BDNF and impaired axonal transport in the striatum. Huntingtin enhances retrograde transport of BDNF, which involves HAP1 and dynactin, but mutant Huntingtin is defective in this process (Gauthier et al. 2004). In Parkinson's disease, there is also reduced axonal transport of BDNF in dopaminergic neurons. The understanding of synthesis, transport, trafficking, and the release of BDNF in diseased conditions will provide useful knowledge for understanding the pathogenesis and underlying molecular mechanism and will help in developing novel therapeutic strategies.

BDNF, a powerful trophic factor with diverse functions, can act as a protector against some cellular damages caused by the neurotoxicity induced by physical and chemical insults. BDNF is a survival factor for cultured rat cerebellar granule neurons and protects them against glutamate-induced neurotoxicity. Restoring BDNF level CEP-1347 reduces mutant Huntingtin-associated neurotoxicity. Overexpression of BDNF is shown to firmly regulate cell survival and differentiation of neuron stem cells (NSC) and protects differentiated NSC against neurotoxicant trimethyltin

(TMT)-induced neurotoxicity through the PI3K/Akt and MAPK signaling pathways. BDNF can also protect against beta-amyloid-induced neurotoxicity both in vitro and in vivo in rats. How BDNF trafficking and release is affected in neurotoxicity would be an interesting area to explore.

#### 7 Conclusion

BDNF is important for neuronal survival and synaptic plasticity. Its synthesis, transport, and release have been intensively investigated since its discovery. It is clear now that BDNF is both anterogradely and retrogradely transported in axonal terminals and it is released in an activity-dependent manner. Although controversies exist regarding the subcellular distribution of BDNF and its exact location of release, more convincing evidence shows that BDNF is stored in presynaptic dense core vesicles, suggesting that BDNF is predominantly transported in an anterograde mode. However, the molecular mechanism underlying intracellular trafficking has not been completely understood, particularly in diseased conditions. Our recent studies provide new evidence that HAP1 and sortilin play an important role in intracellular BDNF trafficking and processing. Future studies should focus on better understanding of more profound molecular mechanisms involved in the trafficking, release, and signaling of BDNF in diseased conditions. Insight into the molecular mechanism of BDNF would help in developing therapeutic treatment for different neurological disorders by targeting mature BDNF and proBDNF.

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# Trafficking of Neurotrophins and Their Receptors and Pathological Significance

Christopher S. von Bartheld

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#### Abstract

To survive and function properly, innervating neurons need to communicate with their targets. This task is accomplished, in part, by trafficking of neurotrophic molecules, in particular the family of nerve growth factor-like molecules, the neurotrophins (NTs). This chapter reviews the intra- and intercellular trafficking of NTs and their receptors with a focus on axonal transport and synaptic transfer, methodology, current concepts, and the relevance for pathological neurological conditions. Neurotrophins are transported by retrograde axonal transport from the neuronal or non-neuronal target cells to the innervating neuronal cell body. The neurotrophic hypothesis postulates that targets produce limiting amounts of trophic substances, including NTs, for

C.S. von Bartheld

e-mail: cvonbartheld@medicine.nevada.edu

Department of Physiology and Cell Biology, University of Nevada School of Medicine, Reno, NV, USA

which innervating neurons compete. NTs bind to NT receptors on axonal terminals and are internalized, presumably in vesicles or endosomes (signaling endosome hypothesis), and are transported by dynein-mediated retrograde axonal transport along microtubules to the cell body where the NT-specific receptors (trkA, B, and C) elicit signal transduction cascades that result in increased neuronal survival, differentiation, and/or plasticity. In addition to the retrograde route, NTs can also be transported anterogradely along axons and can be released at the axon terminal to influence postsynaptic cells. Thus, NTs can transfer in both directions across the synaptic cleft. NT receptors are expressed on the presynaptic as well as postsynaptic membrane, allowing for the activation of signal cascades at pre- as well as postsynaptic sites. NTs such as BDNF and NT-3 are particularly important messengers in synaptic plasticity. In addition to mature NTs, uncleaved precursors (pro-NTs) can also be trafficked and released, and their actions can oppose the functions of mature NTs by activating death signaling through the common NT receptor, p75. Several neurological disorders are currently being investigated for defects in NT trafficking, including Alzheimer's disease, Huntington's disease, Parkinson's disease, neurodevelopmental disorders, epilepsy, glaucoma, hyperalgesia, neuropathies, depression, and eating disorders. Progress in the understanding of physiological and pathological mechanisms of NT trafficking therefore may lead to improved therapies for neurological diseases.

Keywords

Axonal transport • Neurotrophin • Neurotrophin receptor • Neurotrophic hypothesis • Neurodegeneration • Synaptic plasticity

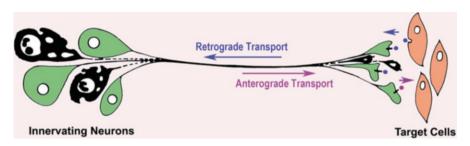
BDNF	Brain-derived neurotrophic factor
CREB	cAMP response element-binding protein
DRG	Dorsal root ganglion
EM	Electron microscopy
ER	Endoplasmic reticulum
GFP	Green fluorescent protein
LDCV	Large dense-core vesicle
LTP	Long-term potentiation
MVB	Multivesicular body
NGF	Nerve growth factor
NT	Neurotrophin
NT-3	Neurotrophin-3
NT-4	Neurotrophin-4
P75	Common neurotrophin receptor
TrkA	Tropomyosin receptor kinase A
TrkB	Tropomyosin receptor kinase B
TrkC	Tropomyosin receptor kinase C

#### 1 Introduction

Intra- and intercellular trafficking of neurotrophic factors is fundamentally important for the neurotrophic hypothesis. This wider framework includes the initial overproduction of neurons during development, followed by programmed (developmental) neuronal death, and the regulation of survival or death of neurons by neurotrophic factors, including the NTs. Besides survival functions, NTs have additional maintenance and plasticity functions in the adult nervous system. Nerve growth factor (NGF) served as the first, prototypical neurotrophic factor. NGF trafficking was closely linked to a paradigm shift in the later half of the 1900s that provided a plausible explanation of how the final number of neurons is matched to the size of their target (systems matching) and how neurons receive feedback from their target. This concept also involves competition for limited trophic supply in the target and trafficking of the NT from the target to deliver the trophic signal to the cell body of the innervating neuron. Early NGF research benefited from some remarkably favorable circumstances that have been reviewed on multiple occasions (Levi-Montalcini 1987; Purves 1988; Oppenheim 1996).

The discovery of the specific and saturable axonal transport of NGF has had a major impact in the field (Hendry et al. 1974; Stoeckel et al. 1975; reviewed from a historical perspective by Oppenheim 1996). Probing neurons with radiolabeled NGF in both in vivo and in vitro settings provided major insights due to high specificity and sensitivity of a quantifiable signal. These studies provided crucial experimental support for the neurotrophic hypothesis. NTs were studied in "Campenot chambers" (Campenot 1977), an experimental approach that allowed to examine NT trafficking in vitro combined with biochemical and molecular analyses (Bhattacharyya et al. 1997; Watson et al. 1999, 2001; MacInnis and Campenot 2002). The 1980s and 1990s saw further seminal progress in NT research with Yves Barde's identification, purification, and sequencing of the second NT, BDNF (Leibrock et al. 1989), and the subsequent expansion of the NT family (NT-3, NT-4). Not only was the entire family of NTs identified but also two types of NT receptors – first p75, then trkA, and other trks (Barbacid 1994; Huang and Reichardt 2003). At the same time, it was realized that signaling and endocytosis are intertwined in important ways (Di Fiore and de Camilli 2001; Sorkin and von Zastrow 2002), and progress was made in the identification of the molecular motors that move organelles and molecules along microtubules: dynein and kinesins and associated motor proteins (Hirokawa 1998; Goldstein and Yang 2000; Vallee et al. 2004). The early 2000s saw the distinction of the roles of mature NTs vs. pro-NTs and increased attention to pro-NT's death signaling capacities (Lee et al. 2001; Lu et al. 2005). The relevance of NTs – especially BDNF – for synaptic plasticity and LTP was realized; this added an entire new dimension in NT function and with that, also new questions and insights into NT trafficking (Poo 2001; Lu et al. 2005).

The scope and focus of this review is the intracellular and intercellular trafficking of NTs and their receptors; the review will consider primarily axonal but also dendritic, synaptic, and somal forms of trafficking. The chapter will also review current knowledge of how NTs are released from neurons and their target cells, how



**Fig. 1** Two major trafficking modes of neurotrophins: retrograde (*blue arrow*) and anterograde axonal transport (*purple arrow*). The neurotrophic hypothesis postulates that developing neurons survive (*green neurons*) only when they successfully compete for target-derived trophic molecules (*blue arrow*) that are internalized at the nerve terminal and are transported retrogradely along the axon (*long blue arrow*) to the cell body. Neurons that fail to obtain a sufficient flow of trophic molecules from the target die by programmed cell death (*black neurons* with condensed chromatin). Neurons also synthesize and send neurotrophins anterogradely along the axon (*long purple arrow*), from the cell body to the nerve terminal, where the neurotrophin is released (*purple*) (Modified and reproduced with permission from the publisher, Elsevier (Barde 1989, *Neuron* **2**, 1525–1534))

they are internalized and transported retrogradely and anterogradely (Fig. 1), and the type of methodology that is used with a discussion of technical challenges and limitations, and finally will examine the relevance of NT trafficking for neurological diseases. For other aspects of NT physiology and pathology, the reader is referred to previous reviews, e.g., on NT history (Oppenheim 1996), NT signaling (Huang and Reichardt 2001, 2003), synaptic plasticity/LTP and synaptic transfer of NTs (Poo 2001; Lessmann et al. 2003), functions of pro- vs. mature NTs (Lee et al. 2001; Lu et al. 2005), internalization of NT receptors in cell lines (Bronfman et al. 2007), retrograde transport of NTs (Reynolds et al. 2000), and retrograde NT signaling (Zweifel et al. 2005).

## 2 Release of Neurotrophins (NTs)

According to the neurotrophic hypothesis, NTs need to be released from target cells that can be neurons or other, non-neuronal cells, such as muscle. NTs can also be released from glial cells. NTs are extremely potent signaling molecules (Thoenen and Barde 1980), and therefore, cells release frustratingly low amounts of endogenous NTs (Gärtner et al. 2000; Dieni et al. 2012, but see Rush 2001, for an alternative view). Low amounts of released NTs are consistent with the neurotrophic hypothesis that postulates that trophic material be released in limiting amounts from target cells and must be competed for by innervating neurons. The apparent low abundance of NTs for release poses a considerable technical challenge, as will be further discussed in Sect. 5. Nevertheless, studies have shown that NTs can be released from different neuronal compartments, including dendrites (Hartmann et al. 2001; Kojima et al. 2001) and axon terminals (von Bartheld et al. 1996a; Fawcett et al. 1997; Kohara et al. 2001; Lever et al. 2001; Wang et al. 2002), while release from cell bodies is less

clear (Blöchl and Thoenen 1995; Balkowiec and Katz 2000). How are NTs or pro-NTs trafficked to sites of release? This is presumably accomplished by trafficking along routes previously established for other peptides, namely, through rough endoplasmic reticulum (ER), transport through the Golgi system, glycosylation, and cleavage of the pro-form in the Golgi system (Conner et al. 1998; Mowla et al. 1999, 2001; Farhadi et al. 2000; Lessmann et al. 2003), while the role of smooth endoplasmic reticulum is less clear (Gärtner et al. 2000). Distinct vesicles containing BDNF can be targeted to either dendritic or axonal compartments (Adachi et al. 2005; Dean et al. 2012).

Cells release peptides via two principal types of release, constitutive and regulated (activity-dependent) modes (Burgess and Kelly 1987). Release of NTs has been reported to be accomplished by both, constitutive and regulated pathways (Blöchl and Thoenen 1995; Goodman et al. 1996; Gärtner and Staiger 2002; Lessmann et al. 2003), although some investigators question the dichotomy and the existence of a true activity-independent baseline of release (Poo 2001). Several groups showed activity-dependent release (Ghosh et al. 1994; Blöchl and Thoenen 1995). Furthermore, the pattern of stimulation frequency is important (Balkowiec and Katz 2000), and release appears to depend on the status of internal calcium stores (Blöchl and Thoenen 1995; Gärtner and Staiger 2002; reviewed by Lessmann et al. 2003). Such release modes are highly relevant for mechanisms of LTP. In this context, properties of a val-met mutation of BDNF are of interest – this mutation does not affect constitutive release, but it impairs regulated BDNF release, apparently due to disruption of a trafficking signal in the BDNF prodomain region (Egan et al. 2003). The resulting defect in BDNF sorting to the regulated secretory pathway has consequences for several neurodegenerative and psychiatric disorders, including depression, eating, and bipolar disorders (Chao et al. 2006), as discussed in Sect. 6 of this review. Which compartments and organelles are involved in NT release? Ultrastructural studies designed to reveal the compartments and organelles involved in NT release from neurites have been inconclusive (Gärtner et al. 2000). Smooth ER membranes were labeled as well as microtubules, but there was no accumulation at synaptic vs. extrasynaptic sites, and large dense-core vesicles (LDCVs) were not labeled. The examination of the ultrastructural distribution in axon terminals has consistently shown NTs to be associated at least to some degree with LDCVs (Michael et al. 1997; Fawcett et al. 1997; Luo et al. 2001; Salio et al. 2007; Dieni et al. 2012). It has been proposed that the apparent lack or paucity of NT immunoreactivity in dendrites (Drake et al. 1999) is due to NTs being "obscured" by trks, so that NT antibodies fail to recognize NTs when they are bound by trk receptors (Rush 2001).

Release at synaptic/extrasynaptic sites – wiring or volume transmission? Being neuropeptides, NTs are supposed to be released primarily at extrasynaptic locations via "volume transmission" (Merighi 2002; Torrealba and Carrasco 2004; Feng et al. 2011), but not directly at the synaptic density. Consistent with such volume transmission, NTs appear to be released from neurites without preference for synaptic sites (Gärtner et al. 2000; Hartmann et al. 2001; Gärtner and Staiger 2002). On the other hand, NTs can be released from axon terminals,

as indicated by studies examining synaptosomes (Androutsellis-Theotokis et al. 1996; Fawcett et al. 1997), DRG terminals (Lever et al. 2001), cultured neuronal networks (Kohara et al. 2001), and retinal ganglion cell axon terminals (Wang et al. 2002). This suggests synaptic release after anterograde axonal transport, but primarily extra- (or iuxta-)synaptic release from postsynaptic sites in dendrites. Recent work indicates that synaptotagmin (Syt-IV) is involved in targeting BDNF-containing vesicles to either dendritic or axonal synaptic sites where they are released by activity-dependent mechanisms (Dean et al. 2012).

Are release mechanisms different between different NTs? Some studies have reported that BDNF release is regulated (or that regulated release occurs over a baseline constitutive component, Blöchl and Thoenen 1995; Lessmann et al. 2003), while NGF (and NT-3) release was reported to always be constitutive (Mowla et al. 1999; Farhadi et al. 2000). However, such differences may be ascribed to differences in experimental protocols rather than true differences between NTs (Griesbeck et al. 1999; Neet and Campenot 2001; Wu et al. 2004), and, as mentioned, some investigators question whether constitutive release truly exists for NTs (Poo 2001).

Initially, only release of mature NTs was envisioned, based on the assumption that all pro-NTs were cleaved intracellularly prior to release (Mowla et al. 1999, 2001; Farhadi et al. 2000). It is now thought that cells differ in the degree of such cleaving, resulting in release of both pro-NTs and mature NTs. If cleavage can occur both intra- and extracellularly, then this provides a novel level of complexity in the regulation of NT availability and function (Lee et al. 2001; Lu et al. 2005). However, because of the low abundance of released endogenous NTs, it is still controversial whether NTs and specifically BDNF are released by neurons in vivo as the pro-NT form or mostly as mature BDNF (Matsumoto et al. 2008; Yang et al. 2009; Greenberg et al. 2009; Dieni et al. 2012). This is an important question that has resulted in an ongoing controversy. Endogenous proBDNF can be transported anterogradely and retrogradely along axons (Wang et al. 2006), but this does not inform about the released form of BDNF. BDNF is easier to measure when overexpressed or tagged, and in vitro, but these conditions may change the release mode, the cleavage properties, and capacity of NT trafficking, and the in vitro condition may not reflect the in vivo situation (see Sect. 5).

Another question is whether the release of newly synthesized NTs is different from that of recycled (internalized) NTs. Based on evidence from other growth factors and examination of exogenous (radiolabeled) NTs, a substantial fraction of released and internalized trophic factors are not immediately degraded, but shuttled into a recycling pathway in neurons (von Bartheld et al. 2001; Rind et al. 2005). Depending on how much NT may be recycled rather than degraded, a major fraction of NTs in neurons may not comprise newly synthesized NT, but rather recycled NT after its initial release and internalization (von Bartheld et al. 2001; Lessmann et al. 2003). When recycled, what is the storage compartment? Multivesicular bodies (MVBs) are strong candidates for NT storage – NT-filled MVBs are often localized adjacent to synapses on the postsynaptic side (Rind et al. 2005; von Bartheld and Altick 2011). It has been shown that the recycled pool of BDNF indeed is relevant for LTP (Santi et al. 2006).

What is the role of NT receptors in release? NT receptors need to be targeted to pre- as well as postsynaptic membranes (Ascaño et al. 2009), indicating that NTs and/or pro-NTs can be released at synapses and may bind to pre- as well as postsynaptic NT receptors. Truncated trkB receptors have a role for BDNF release from glial cells (Alderson et al. 2000) and possibly neurons as well (Biffo et al. 1995).

#### 3 Retrograde Axonal Transport

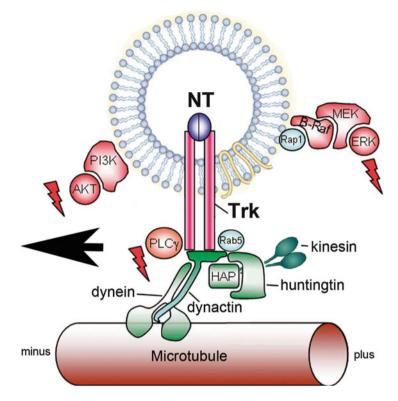
This section will follow the pathway of NTs when they are internalized by axon terminals and sent onto a trafficking route along the axon (a distance that may exceed 1-2 m) and that culminates with the delivery of the NT (and NT-induced signals) to the neuronal cell body. The evidence for receptors and organelles involved in this process will be reviewed, and the significance of the prevailing "endosome signaling hypothesis" and its alternatives discussed.

Internalization. It has been appreciated for a long time that the internalization of NTs is receptor mediated (Hendry et al. 1974; Stoeckel et al. 1975; Reynolds et al. 2000). The NT-receptor complex can be endocytosed by either clathrin-mediated or nonclathrin-mediated pathways (Howe and Mobley 2004; Bronfman et al. 2007). The socalled caveosomes may concentrate lipid-raft-derived NT signals (reviewed in Howe and Mobley 2004). Thus, in order to achieve internalization of NTs at the axon terminus, NT receptors need to be targeted first to, and incorporated into, the axon terminal plasma membrane. Which NT receptors mediate internalization and retrograde axonal transport of NTs? This has been examined extensively in the 1990s, and various lines of evidence indicate that NGF is internalized and transported in neurons primarily by trkA, while for other NTs (BDNF, NT-3 and NT-4) both trk receptors and p75 receptors contribute, and NT-4 appears to rely the most on p75 as a transporter (Curtis et al. 1995; von Bartheld et al. 1996b). p75 is believed to be expressed in vivo in a ratio of about fiveto-tenfold higher than trk receptors (Chao 1994), and the relationship between p75 and trk receptors and the functional significance of co-expression of these two receptor types has been intensely studied (Bothwell 1995; Bronfman et al. 2007). Co-expression of p75 and trk receptors appears to create a higher-affinity binding site for NTs (Chao 1994; Bothwell 1995; Esposito et al. 2001). Initial models suggested a simultaneous binding of p75, trk, and the NT in a trimeric complex, but this is not consistent with more recent molecular modeling data (Wehrman et al. 2007; Barker 2007). It is possible that p75 receptors "present" the NT to the less common trk receptors (Biffo et al. 1995; Barker 2007). It has been suggested that the NT is initially captured by the more abundant p75 receptors and over time (during axonal transport) accumulates on trk receptors in the signaling endosome (Butowt and von Bartheld 2009; Chowdary et al. 2012). This proposed mechanism is consistent with the known differences in on-rates and off-rates (the p75 receptor has a faster off-rate than the trk receptors – Rodriguez-Tébar and Barde 1988; Bothwell 1995). NT-containing endosomes have been reported to co-localize p75 and trk receptors to varying degrees, and some studies suggest that p75-containing endosomes are routed differently than trk-containing endosomes (Hibbert et al. 2006; Bronfman et al. 2007). It has been known for a long time that

NTs injected directly into the cell body have no trophic effect, demonstrating that the binding of NTs to the cell surface NT receptor and the internalization of the ligand-receptor complex are essential for mediating the trophic signal (Claude et al. 1982; critically reviewed by Neet and Campenot 2001). Several molecules involved in the internalization and transport of NTs and NT receptors have been identified; these include pincher and Rabs (Rab5 and Rab7) (Heerssen et al. 2004; Saxena et al. 2005; Valdez et al. 2005; Deinhardt et al. 2006; Bronfman et al. 2007). The p75 receptor can be cleaved and the C-terminal p75 fragment may be trafficked; this fragment has been shown to retain signaling capacity (Kanning et al. 2003). In summary, there is abundant evidence for retrograde axonal transport of NTs (reviewed in Reynolds et al. 2000) as well as the NT receptors, p75 and trks (Johnson et al. 1987; Ross et al. 1994; Ehlers et al. 1995; Mufson et al. 1999; Moises et al. 2007).

The endosome signaling hypothesis and two alternatives. Evidence for the receptor-mediated internalization of NTs and emerging knowledge about endocytosis of growth factor receptors and their ligands prompted the so-called "endosome signaling" concept: Pinching off NT receptor containing plasma membrane results in vesicles with the NT bound on the inside and the tyrosine kinase domain of the NT receptor on the outside of the vesicle, available for docking of additional signaling molecules (Fig. 2). A large number of studies have accumulated supporting evidence for the validity of this model: Phosphorylated (activated) trkA is retrogradely transported (Ehlers et al. 1995; Bhattacharyya et al. 1997); signaling endosomes can be purified from nerves (Delcroix et al. 2003); they contain the relevant signaling molecules depicted in Fig. 2; some of the NT receptors possess moieties that directly attach to the molecular motor (dynein) that moves the endosome retrogradely along microtubules (Yano et al. 2001; Yano and Chao 2004); these vesicles retain signaling capacity, as reviewed by multiple groups (Zweifel et al. 2005; Wu et al. 2009; Oppenheim and von Bartheld 2008). Some of the signaling may occur already during transit, since sufficient signal transducers are attached to the NT- and NT-receptor-containing vesicle. However, most of the signaling is believed to occur once the signaling vesicle is delivered to the cell body. At the cell body, signaling cascades (via CREB and other pathways) regulate gene expression in the nucleus (Riccio et al. 1997); after delivery of the signal, the signaling endosome is believed to be dismantled, possibly by transition to a MVB and eventual to a lysosome (Mukherjee et al. 1997; von Bartheld and Altick 2011). Alternatively, the endosome or components of the signaling endosome are recycled by targeting to other cellular compartments (e.g., synapses on dendrites, von Bartheld et al. 2001; Rind et al. 2005). Currently, there is widespread support among investigators in the field for the signaling endosome hypothesis.

One alternative to the signaling endosome hypothesis is the concept that NTs may signal locally within the axon terminus and that this generates one or more second messengers in the axon terminal that is moved retrogradely along the axon. There is some evidence that second messengers other than axonally transported NTs or NT receptors contribute to the retrograde signal; these include G proteins as well as MEK kinase, MAP kinases, and PI3 kinases and/or their subunits



**Fig. 2** The signaling endosome. The internalized neurotrophin (*NT*)-receptor (*Trk*) complex is transported retrogradely from the nerve terminal to the cell body. The internalized vesicle serves as a platform for docking of proteins that elicit signal transduction cascades (*red*): via the phospholipase C-gamma (*PLC* $\gamma$ ), Raf-MAPK-ERK, and PI3K/AKT signaling pathways (*red lightning symbol*); molecular motors and regulators (*green*) attach for transport along microtubules (*brown*). The *arrow* indicates the direction of transport toward the minus end of microtubules. For details and references, see Zweifel et al. 2005 (*Nature Reviews Neuroscience* **6**, 615–625) and Gauthier et al. 2004 (*Cell* **118**, 127–138) (Material adapted, with permission, from: Oppenheim and von Bartheld 2008 (Fundamental Neuroscience, 2nd edition, Academic Press))

(Johanson et al. 1996; Reynolds et al. 2000; Weible and Hendry 2004). It is unclear to what extent the transported NT-containing vesicle or endosome may contribute to the signaling beyond the initial signaling at the axon terminus; in the strongest form of the second messenger concept, the retrograde axonal transport of the NT may simply serve to remove the NT from the region of the axon terminus ("vacuum cleaner model", Bothwell 1995), so that the local NT signal can be terminated and does not continue to elicit signals.

A third possibility is the "wave model," which is based on the surprising observation that at least in some cases, a trophic signal arrives at the level of the cell body prior to the time needed for the NT-containing endosome to arrive there (Senger and Campenot 1997) and that consequently some signaling occurs that is independent of the retrograde axonal transport of the NT-containing vesicle that

initiated the signal (Miller and Kaplan 2001; Neet and Campenot 2001; MacInnis and Campenot 2002). The major weakness of this model is that there is no compelling physical mechanism of how the signal could spread as a lateral "wave" along the required distances (Ginty and Segal 2002; Howe and Mobley 2004; Wu et al. 2009). Furthermore, current evidence is based on a small set of in vitro experiments, and some assumptions are made about identical properties of radiolabeled vs. native NTs.

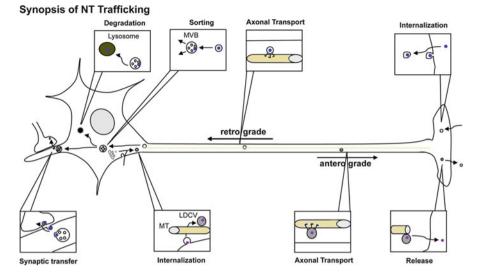
Transport organelle. Considering the long distance from the axon terminus to the cell body, it is surprising that there is very little direct evidence for the type of organelle that carries the NT signal retrogradely along the axon. Two major models can be distinguished: The signaling organelle is a multivesicular body (MVB, Parton et al. 1992; Valdez et al. 2005; Weible and Hendry 2004; Philippidou et al. 2011), or it is a small vesicle or endosome (Delcroix et al. 2003; Altick et al. 2009: Wu et al. 2009: von Bartheld and Altick 2011). The former notion is based primarily on in vitro experiments performed with large tags of the NT, which recent work has shown to generate abnormal organelles or quantities of organelles (von Bartheld and Altick 2011). The alternative view is that the endosome is a small vesicle and remains such a structure up to the level of the cell body. Purification experiments support this view (Delcroix et al. 2003), as do experiments with quantum dot-labeled NTs (Cui et al. 2007). MVBs were ruled out as the "normal" carrier based on their low frequency in normal nerves in vivo and lack of labeling when the ultrastructural distribution of radiolabeled NTs was determined within nerves (Altick et al. 2009). While there is no doubt that retrogradely transported NTs eventually accumulate in MVBs within the cell body (Claude et al. 1982; Rind et al. 2005; von Bartheld and Altick 2011), NTs do not appear to travel within this type of organelle while en route within the axon in vivo.

Arrival of the NT signal in cell body. The signal transduction pathways activated by NTs in the cell body have been reviewed elsewhere and will not be repeated here (Huang and Reichardt 2001, 2003). There has been one major aspect that has puzzled the investigators: How is it that the signal within the signaling endosome is so long-lived? When the same signaling cascades are activated by other ligands and other types of tyrosine kinase receptors, the MAP kinase activation is measured within minutes, not hours (Kao et al. 2001; Wu et al. 2001). Prolonged activation appears essential, since it can take hours to move a signaling endosome (or the NT) from the axon terminus to the cell body, e.g., in the motor neurons of the spinal cord. Several solutions have been offered why NT binding to trks in neurons may induce longer-lasting signal duration (Kao et al. 2001; Wu et al. 2001; Saxena et al. 2005). The concentration of trk receptors in the endosome is sufficiently high to dimerize and autophosphorylate (Miller and Kaplan 2001). Another mechanism in play could be that the NT comes off one NT receptor and binds another NT receptor within the same vesicle (Butowt and von Bartheld 2009; Chowdary et al. 2012). Does the cell body somehow know where the trophic signal was generated (axon target vs. dendritic afferents)? This question was examined by Segal and her colleagues, and it appears that some signal transduction pathways (ERK) have certain specificity for where the NT signal was generated (Watson et al. 2001; Heerssen and Segal 2002). However, another study found that the same ERK was contained in signaling endosomes (Delcroix et al. 2003), indicating that the type of ERK activation may be variable between different cell types or other conditions (Bronfman et al. 2007).

Fate of NTs after arrival in the cell body. How is the NT signal "turned off"? One solution that has been proposed is that the NT endosome is incorporated into the internal vesicles of a MVB, essentially preventing access of the NT receptor to the cytosol (Sorkin and von Zastrow 2002; von Bartheld and Altick 2011). Initially, it was assumed that NTs are degraded immediately or within a few hours after signaling (Claude et al. 1982; Reynolds et al. 2000; Neet and Campenot 2001). Some neurotrophic factors such as cardiotrophin-1 indeed appear to be rapidly degraded in lysosomes (Rind et al. 2005). However, other trophic molecules are used multiple times prior to degradation, and NTs appear to belong to this group (von Bartheld et al. 2001; Rind et al. 2005; Sharma et al. 2010). Thus, there are two pools of NTs: newly synthesized NTs and internalized NTs that can be recycled. In fact, recycled, internalized NTs have been shown to serve important functions in LTP (Santi et al. 2006). One of the molecular mechanisms that decides between degradation and recycling of retrogradely transported NTs appears to involve phosphorylation of the p75 receptor (Butowt and von Bartheld 2009). While there has been progress in elucidating some of the mechanisms of retrograde neurotrophin signaling, major gaps in understanding of the transport and arrival of the signaling complex remain to be filled. The major steps of retrograde and anterograde trafficking of NTs are summarized in the schematic of Fig. 3.

# 4 Anterograde Axonal Transport

Based on the neurotrophic hypothesis and the early discovery of retrograde axonal transport of NGF and other NTs (Hendry et al. 1974; DiStefano et al. 1992), trafficking of NTs was for several decades thought to occur primarily if not exclusively in the direction toward the cell body. Despite abundant evidence for trophic effects of afferents on neurons (Linden 1994), it came as a surprise when it was discovered in 1996 that both exogenous and endogenous NTs can move anterogradely along axons (von Bartheld et al. 1996a; Zhou and Rush 1996) and that NTs can be transferred from presynaptic to postsynaptic sites (von Bartheld et al. 1996a). These first reports were rapidly confirmed (Altar et al. 1997; Johnson et al. 1997; Altar and DiStefano 1998; Conner et al. 1998). The discovery of anterograde axonal transport of NTs opened a new venue of NT research: Can NTs behave like neurotransmitters? Are they released at the axon terminus by activity-dependent mechanisms? In which organelles are they stored there, and how are they released? What are the functions of this new trophic pathway?



**Fig. 3** Synopsis of retrograde and anterograde trafficking of neurotrophins (*NTs*). The upper half of the schematic illustrates the major steps of retrograde traffic, from internalization at the axon terminus, via microtubule-mediated axonal transport along the axon, sorting in the cell body and eventual degradation in lysosomes. The lower half of the schematic shows the major steps of synthesis or internalization of NTs in the cell body, microtubule-mediated anterograde axonal transport, and release at the axon terminus. In addition, synaptic transfer of NTs via MVBs is indicated. Abbreviations: *LDCV* large dense-core vesicle, *MT* microtubule, *MVB* multivesicular body

Some investigators now believe that anterograde axonal transport of NTs is quantitatively more significant than the "conventional" retrograde axonal transport (Conner et al. 1998; Adachi et al. 2005). Despite its moment of fame in the late 1990s, anterograde axonal transport of NTs has been studied much less than retrograde transport, and therefore, less is known about its mechanism. Anterogradely transported NTs are now believed to be mostly stored within LDCVs in axon terminals (Michael et al. 1997; Fawcett et al. 1997; Luo et al. 2001; Salio et al. 2007; Dieni et al. 2012), although some controversy remains. The amounts of both endogenous and exogenous NTs that are transported anterogradely appear to be lower, on a per neuron basis, than those that can be transported retrogradely (von Bartheld et al. 1996b). Anterogradely transported neurotrophins have been shown to have multiple effects on their afferents, not only with regard to neuronal survival but also synaptic transmission and dendritic morphology (Caleo et al. 2003; Caleo and Cenni 2004; von Bartheld 2004).

It should be noted that anterograde axonal transport of NTs and NT receptors serves two very different functions: NT anterograde transport appears to mediate trophic signals that arise from afferents (Linden 1994), as opposed to the well-established roles of efferent targets, while anterograde transport of NT receptors primarily delivers the NT receptors that are needed in the axon terminus for

internalization and retrograde transport of target-derived trophic signals (Ascaño et al. 2009). While it is known that the anterograde axonal transport of NTs in adults does not require p75 receptors, the role of trk receptors is less clear, although they seem to contribute to anterograde axonal transport of BDNF (Butowt and von Bartheld 2005). Anterograde axonal transport of NTs is mediated by conventional kinesins (Butowt and von Bartheld 2007), while anterograde axonal transport of trk receptors is facilitated by sortilin (Vaegter et al. 2011), and the association between trkB receptors and kinesin-1 is accomplished by the protein JIP3 (Huang et al. 2011).

Release mechanisms of anterogradely transported NTs have to be studied specifically at the axon terminus, which is not an easy task. Such release was first examined in synaptosomal preparations (Androutsellis-Theotokis et al. 1996; Fawcett et al. 1997), and subsequently by utilizing advantageous model systems such as the DRG central projection into the spinal cord (Lever et al. 2001), retinal ganglion cell axons in the optic tectum (Wang et al. 2002). In addition, one can culture neurons and let them form synaptic connections, engineered to express BDNF-GFP in the presynaptic neuron and observe labeled puncta to transfer from presynaptic to postsynaptic sites (Kohara et al. 2001). These studies showed that such pre- to postsynaptic transfer of NTs was activity dependent and modulated by numerous compounds, including intracellular calcium, cAMP, NMDA receptors, and stimulation pattern (Kohara et al. 2001; Lever et al. 2001; Wang et al. 2002).

Anterograde axonal transport of internalized NTs. While much of the anterograde axonal transport of NTs presumably involves newly synthesized NTs, exogenous NTs can also be targeted to an anterograde axonal pathway (Butowt and von Bartheld 2001). Such exogenous NTs are internalized by retinal ganglion cells, traverse the Golgi system, and thereby join what is presumably the "normal" anterograde axonal route. Some of the NTs in this pathway are packaged in LDCVs (Wang et al. 2002), and they are released from axon terminals in the optic tectum in an activity-dependent fashion (Wang et al. 2002). One exogenous NT, NGF, was shown to be immunoprecipitated by p75 receptor antibodies in this pathway to the optic tectum where the same NGF subsequently (after release) binds primarily to trkA receptors and embarks on a new route to the locus ceruleus (Feng et al. 2011). Thus, NT itineraries can be linked in neural networks in different anterograde and retrograde trafficking combinations (Figs. 1 and 3).

## 5 Methodology

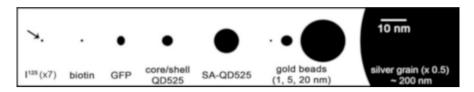
This section reviews the types of techniques that researchers use to investigate trafficking of NTs. It will examine limitations and pitfalls, as well as strategies to overcome challenges, and discuss some current controversies in the field.

When endogenous NTs are examined, they can be visualized with NT-specific antibodies (Conner 2001). Another approach is to express a specific tag (such as GFP or a myc tag) with the NT of interest. Obviously, all NTs recognized by the antibody or the tag signal itself will be visualized (as in a snap shot), and therefore,

this approach alone does not reveal the trafficking itself (either qualitative or quantitative). For example, the presence of the NT in the axon suggests axonal transport, but it does not indicate the direction (anterograde or retrograde) or the velocity and capacity. Some hints may be given by the lack of synthesis (by in situ hybridization) of the NT in the cell body or the target cells (Conner et al. 1997), but if expression levels are very low, no firm conclusions can be made about trafficking. A useful and common technique is to apply a ligature to a nerve and to determine whether the NT accumulates distally or proximally to the ligature or both (Johnson et al. 1987; Ross et al. 1994; Ehlers et al. 1995; Mufson et al. 1999). This gives information about the direction of axonal transport and a rough estimate of the relative amount. Caveats are that glial cells surrounding the site of ligature may be induced to express NTs, and they may thus accumulate in response to such nerve injury, resulting in false-positives.

A major problem with the study of endogenous NTs is the extremely low abundance of the endogenous NTs in vivo. This is a challenge, because the methods used to visualize the NT may not have sufficient sensitivity. In addition, NTs may simply be present in concentrations too low to be detected, yet high enough to have significant trophic effects. Therefore, lack of label cannot be interpreted as evidence for a lack of NTs. Furthermore, based on the localization of NTs, antibodies may fail to reveal the NT, for example, if the antibody does not recognize the NT when it is bound to certain receptors (Rush 2001). One solution to the problem of low abundance is to overexpress the NT of interest. However, this approach may saturate normal trafficking capacities and thus lead to false conclusions about trafficking. Another approach is to transfect tissues or cells with a molecular tag on the NT – this, however, may change NT trafficking, receptor binding, sorting and degradation, as well as kinetics. This is especially important to consider when the tag is large relative to the size of the NT protein (Möller et al. 1998; Parak et al. 2005; Chowdary et al. 2012).

When exogenous NTs are examined, some of the abovementioned problems (such as not knowing the direction and quantity of trafficking) are minimized when the NT is introduced into a well-defined compartment. However, the exogenous NT has to be different or tagged so that it can be distinguished from the endogenous NT already present in the system. There are several different options of how NTs can be tagged (Fig. 4). Particularly important, historically, has been the tagging of NTs with the isotope 125I (radioiodination). This procedure only minimally modifies the NT, so the radiolabeled NT's binding properties are unaffected and the NT maintains bioactivity (Johnson et al. 1978; Vale and Shooter 1985; Rosenfeld et al. 1993; von Bartheld 2001). The probe is highly sensitive and quantifiable and can be localized at the ultrastructural level without compromising tissue quality, since probe penetration through tissue is not an issue as it is with antibody visualization. Drawbacks are that the probe cannot be tracked in real time and that the resolution at the EM level is poor, allowing only localization of the NT (silver grain) with major organelles and large enough structures such as synapses (Claude et al. 1982; von Bartheld 2001; Rind et al. 2005). Other options for tagging NTs include biotinylation (Rosenberg et al. 1986), conjugation with GFP (Gerdes and



**Fig. 4** Comparison of the sizes of commonly used protein tags. I125 is the smallest tag (seven times smaller than indicated). The core/shell of quantum dots (QDs) is similar in size to green fluorescent protein (*GFP*). Gold particles come in various sizes, as indicated. After autoradiographic development of radiolabel in monolayer emulsion, silver grains are the largest with diameters of 100–200 nm, thereby relinquishing the small size of the radiolabeled tag. Abbreviation: *SA* streptavidin

Kaether 1996; Haubensak et al. 1998; Watson et al. 1999; Kohara et al. 2001; Adachi et al. 2005; Guillemot et al. 2007), myc tagging (Möller et al. 1998), gold particle tagging (Sandow et al. 2000; Salio et al. 2007), and conjugation with quantum dots (Giepmans et al. 2005; Vu et al. 2005; Cui et al. 2007; Rajan et al. 2008; Altick et al. 2009; Chowdary et al. 2012; Xie et al. 2012). With the exception of biotinylation and myc tagging, these procedures result in a NT-tag complex that is considerably larger than the original, native NT, and the obvious concern is that such a protein modification may change trafficking properties of NTs. Typical controls include verification that the tagged NT still binds to its receptors, has bioactivity, and that it is internalized (Vu et al. 2005; Cui et al. 2007; Rajan et al. 2008; Xie et al. 2012). However, investigators rarely test tagged NTs for any trafficking changes, including capacity, speed, sorting, intracellular aggregation, organelle transition, and degradation kinetics (Neet and Campenot 2001). Indeed, it is not known how such tags affect the NT itineraries, sorting, recycling, and degradation and whether and at what point the tag may dissociate from its target protein (Tekle et al. 2008; Wiggins et al. 2012).

Another problem with expression of either tagged NTs or overexpression of tagged or native NTs is that this may result in much higher levels of the NT than normal, and this may saturate the normal capacity of NT processing, cleavage, trafficking, sorting, and release and accordingly may lead to false-positives, because the tagged and/or overexpressed NT may partially or entirely be directed to abnormal trafficking routes and sites of accumulation (Hartmann et al. 2012; Wiggins et al. 2012). Some of these issues may explain discrepant findings and current controversies, for example, the question whether neurons release BDNF mostly as a mature BDNF form or whether also the uncleaved proBDNF is released by neurons in vivo (Matsumoto et al. 2008; Yang et al. 2009; Greenberg et al. 2009; Dieni et al. 2012). In this context, it is important to remind the reader that caution is also advised when experiments are performed in vitro as opposed to in vivo. For example, extracellular BDNF is rapidly cleaved in vivo by glial cells; when neurons are maintained in an artificial environment lacking glial cells, extracellular proBDNF can persist for a longer time (Yang et al. 2009). Also, neurites (neuronal processes) grown in vitro tend to express many more MVBs than they would in vivo, and application of exogenous NTs conjugated to particles such as gold can induce formation of multivesicular bodies and incorporation of NTs into such organelles, when this would not normally happen (von Bartheld and Altick 2011). Nevertheless, trafficking studies in Campenot chambers and microfluidic chambers have considerably extended knowledge and concepts of NT trafficking (MacInnis and Campenot 2002; Cohen et al. 2011). Examination of NT receptors rather than NT ligands is technically more robust, since receptors are expressed at higher levels than NTs and therefore are easier to visualize, both by immunological and GFP-expression methods. Furthermore, NT receptors stay confined to the producing cells and typically are not released from cells and do not transfer from one cell to another cell (with the possible exception of transfer via exosomes, von Bartheld and Altick 2011). Taken together, results obtained from in vitro studies, and results obtained by observing trafficking of grossly-modified NTs, need to be interpreted with considerable caution.

## 6 Diseases and Disorders Related to NT Trafficking

With the emergence of the neurotrophic hypothesis, it was suggested that the neurotrophic concept may not only apply to the development and differentiation of neurons but also to their maintenance. Accordingly, neurodegeneration may be the result of a failed supply of neurotrophic signals (Appel 1981; Saper et al. 1987). Based on this concept, neurodegenerative diseases may result from a reduced production, reduced axonal transport, or reduced efficacy of neurotrophic signaling. Many neurodegenerative diseases manifest with axonal transport problems (Krüttgen et al. 2003; Bronfman et al. 2007; Perlson et al. 2010; de Vos et al. 2008), and some neurological disorders have been associated with specific NTs (Shu and Mendell 1999; Bender 2004; Chao et al. 2006; Perlson et al. 2010; Hartmann et al. 2012). Abnormalities in neurotrophin signaling during formation of neural circuits may lead to disorganization of those circuits and predispose to neuropsychiatric disease during later life (Angelucci et al. 2005; Autry and Monteggia 2012). This section reviews current evidence for defects or abnormalities in NT or NT receptor trafficking for several key neurological disorders, including Alzheimer's disease, Huntington's disease, Parkinson's disease, Rett and fragile X syndromes, epilepsy, glaucoma, hyperalgesia, neuropathies, as well as psychiatric disorders such as schizophrenia and eating disorders.

## 6.1 Neurodegenerative Diseases

*Alzheimer's disease*. The hallmark of Alzheimer's disease is the degeneration of cholinergic basal forebrain neurons, among other cell types. These neurons are NGF dependent, and mouse models of Alzheimer's disease suggest that abnormalities in the retrograde axonal transport of NGF play a pathophysiological role (Cooper et al. 2001; Salehi et al. 2009). Increased levels of amyloid precursor protein (APP) appear to cause a specific NGF trafficking defect in cholinergic

basal forebrains in a mouse model (Salehi et al. 2006). Enhancing levels of NGF has been tested as a therapeutic approach in patients with early stage Alzheimer's and showed potential effectiveness (Tuszynski et al. 2005), but this phase 1 clinical trial also revealed some of the challenges in NT delivery to the brain (Tuszynski 2007).

*Huntington's disease*. In the heritable Huntington's disease, neurons degenerate in the basal ganglia, primarily the caudate/putamen. The key mutation of this disease is huntingtin, a protein that attaches to the molecular motor that moves cargo along microtubules, among other functions (Gauthier et al. 2004; Ross and Tabrizi 2011). Cortical neurons produce BDNF and transport it and other cargo molecules by anterograde axonal transport to the striatum (Charrin et al. 2005), but when huntingtin is mutated, the anterograde trafficking of BDNF to striatal synapses is abnormally reduced. Indeed, supply of exogenous BDNF can improve symptoms of Huntington's in animal models of the disease (Zuccato and Cattaneo 2007). Accordingly, this is a prime example of the physiological significance of a trafficked NT for disease progression and therapeutic improvement using targeted NTs. In other motor degenerative disease such as amyotrophic lateral sclerosis, trafficking of neurotrophic factors does not appear to be central to disease pathology (Marinkovic et al. 2012; Redler and Dokholyan 2012).

*Parkinson's disease*. The cause of Parkinson's disease is the degeneration of dopaminergic neurons in the substantia nigra, pars compacta. It is suspected that neurotrophic factors are involved, including NTs, but the precise cause and problem (expression, transport, receptors), susceptibility to toxic factors and cytokines, reduced levels of neurotrophins, and thus potential therapeutic strategies are not yet clear (Siegel and Chauhan 2000; Rangasamy et al. 2010).

# 6.2 Neurodevelopmental Disorders

Rett syndrome is a severe neurodevelopmental disorder with regressive cognitive and language skills, caused by a mutation in the MeCP gene. Brains of patients with Rett syndrome have reduced dendritic growth and spine densities in cortical neurons. Not only BDNF expression, but possibly also BDNF trafficking and secretion are affected (Zeev et al. 2009). Fragile X syndrome is another neurodevelopmental disorder, with impairment of BDNF vesicle docking and fusion (densecore vesicle docking). Thus in both syndromes, BDNF expression, trafficking, and/ or release are suspected to play key roles in the disease process (reviewed in Hartmann et al. 2012).

### 6.3 Other Disorders

*Glaucoma*. Glaucoma is characterized by progressive optic nerve degeneration and retinal ganglion cell death due to increased intraocular pressure. It has been shown that a major factor causing retinal ganglion cell death is the reduction in BDNF

retrograde axonal transport along the optic nerve fibers from central targets (Quigley et al. 2000; Almasieh et al. 2012). Thus, the sustained retrograde axonal trafficking of an NT is a major determinant of neuronal survival in the adult nervous system.

*Epilepsy*. Seizures induce BDNF mRNA upregulation that leads to increased BDNF production by the dentate granule cells, enhanced anterograde transport and release of BDNF from mossy fiber axons, and activation of either pre- or postsynaptic trkB receptors in the hilus and CA3 stratum lucidum (reviewed in Bender 2004).

*Hyperalgesia and Charcot-Marie-Tooth neuropathy.* The anterograde axonal transport of BDNF in DRG neurons affects pain thresholds (Shu and Mendell 1999). Several proteins controlling endosomal traffic have recently been shown to be affected in peripheral neuropathies. The small GTPase Rab7 regulates trkA trafficking (Saxena et al. 2005). Dynamin and Rabs affect the traffic of several different types of endosomes, but do not appear to be specific for endosomes containing NTs (Bronfman et al. 2007). Therefore, these traffic regulatory proteins likely affect multiple signaling systems.

*Psychiatric diseases.* Bipolar disorder, schizophrenia, eating disorders, and depression have been associated with BDNF (Angelucci et al. 2005). The BDNF hypothesis of neuropsychiatric diseases postulates that BDNF is required developmentally for normal circuit formation and also in the mature CNS to maintain plasticity and function. Diseases specifically linked to BDNF included eating disorder, depression, bipolar disorder, and schizophrenia (reviewed in Chao et al. 2006). A mouse model with a BDNF mutant (Egan et al. 2003) suggested that a relatively common BDNF mutation in humans causes anxiety and depression-like behavior (Chen et al. 2006). However, this simple hypothesis that a traffic-impaired BDNF mutant caused depression or anxiety did not hold up with further scrutiny and meta-analyses (Groves 2007; Frustaci et al. 2008; Verhagen et al. 2010). This does not call into question a fundamentally important role of BDNF in the pathophysiology and/or treatment of these diseases. BDNF's role appears well supported for eating disorders and bipolar disorder/schizophrenia, but the precise roles of BDNF need further clarification.

# 7 Conclusion

Trafficking of NTs is important in neuronal development, for neuronal plasticity, and in the adult nervous system. Ideal tagging of NTs is still hypothetical, and limitations of current tagging methods need to be taken into account. It has major consequences that NT trafficking can be altered by impairments of the molecular motors that move NTs along axons and in endosome traffic. Although the details of the pathophysiological mechanisms are still to be worked out, there is hope that therapeutic interventions can be designed to remove the toxic insult, halt the degenerative process, and preserve the remaining neurons in neurodegenerative disorders.

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Part V

Diseases and Disorders Relevant to Neurotoxins

# Drug Abuse Neurotoxicity: Alcohol and Nicotine as Developmental Stressors

# Trevor Archer, Serafino Ricci, and Max Rapp-Ricciardi

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#### Abstract

Drugs of abuse have the property of inducing adverse health complications, not least neurotoxicity under conditions where both the environmental conditions and activity states associated with their intake may strongly enhance drug toxicity, thereby causing life-threatening health complications and tragedy for relations and caregivers. While both chronic alcohol and/or nicotine abuse induce a variety of neuropathological effects, including damage to the brain, the extent of damage and disruption observed in the developing brain and CNS is a considerable affliction for the affected individuals. On the basis of laboratory and clinical studies, the potential of chemicals, including therapeutic and abused agents, to induce neurotoxic effects has been assessed, with considerations of abuse drugs neurotoxicity encompassing several factors that may accelerate and complicate prevailing conditions: the type and influence of environmental

T. Archer (⊠) • M. Rapp-Ricciardi Department of Psychology, University of Gothenburg, Gothenburg, Sweden e-mail: trevor.archer@psy.gu.se

S. Ricci

Department of Anatomy, Histology, Forensic Medicine and Orthopaedics, Sapienza University of Rome, Rome, Italy

conditions, the presence of daily habits such as coffee breaks/smoking breaks, nutritional status, and neuroimmune system mobilization. Abuse neurotoxicity at several stages of early development, alcohol neurotoxicity, nicotine neurotoxicity, and combinations of alcohol-nicotine neurotoxicity present a threatening scenario of two compounds, benefitting from legality and availability that nevertheless have such potential for destruction over multiple domains, particularly in the undeveloped brain.

Keywords

Abuse drugs • Alcohol • Apoptosis • Development • Nicotine • Oxidative stress • Smoking

## 1 Introduction

The potential of chemicals, including therapeutic and abused agents, to induce neurotoxic effects may be assessed at (i) early developmental, (ii) adolescent/ preadolescent, and (iii) adult stages of individuals' life cycle. In each case, the level of neurotoxicity or neurotoxic propensity remains a singularly important regulatory drug issue and a uniquely significant area of public concern, not least due to the preponderance of agents that have been characterized or are still to be characterized (Judson et al. 2009) due to logistical constraints (OECD 2007). Normal brain development follows a preprogrammed pathway, whether at neuronal, circuitry, or regional levels, that differentiates specific and distinctive regional structural characteristics, phenotypes, and functions in the infant, adolescent, and adult (Bayer et al. 1993; Herschkowitz et al. 1997). The influence of epigenetic factors (e.g., mutant strains), of evermore critical importance (Dufault et al. 2012; Marti et al. 2007), reflects the sensitivity and responsiveness of human and animal brains in constantly changing circumstances regulating gene expression profiles (Archer et al. 2011). The course of brain development may be altered by a wide range of exogenous agents thereby threatening the integrity of affected brains with permanent consequences (Stanwood and Levitt 2004) that culminate in dire neurodevelopmental (Llop et al. 2012) and neuropathophysiological (Miller et al. 2012) outcomes. The enormous range of exogenous agents capable of disrupting the integrity of the affected brain, within defined parameters and conditions, may wreak damage that is all too often permanent and irreversible (Archer 2011). In order to confront the neurotoxic propensities of abused drugs, alcohol and nicotine (smoking), and their combination, are chosen as "drug cases" for several reasons, not least the situation that both are "legal drugs" judicially in most countries. Alcohol presents the greatest social and monetary chemical risk to individuals' well-being and health, and alcoholism is a multifactorial, genetically associated disorder with major negative issues, highly frequent and a strong cause of premature death.

Considerations of abuse drugs neurotoxicity must take into account several factors that may accelerate and complicate prevailing conditions: the type and influence of environmental conditions, the presence of daily habits such as coffee

breaks/smoking breaks, nutritional status, and neuroimmune system mobilization. It ought to be borne in mind that the environmental conditions and activity states associated with their intake are likely to enhance markedly drug toxicity levels, thereby developing a situation wherein continued abuse leads not only to adverse health complications, including neurotoxicity, but also to life-threatening health complications. For example, animal experiments indicate that even moderate increases in environmental temperatures and physiological activation, the conditions typical of the so-called "rave parties," dramatically potentiate brain hyperthermic effects of psychomotor stimulants, particularly methamphetamine (METH) and 3,4-methylenedioxymethamphetamine (MDMA). Kiyatkin and Sharma (2012) have demonstrated that methamphetamine also induces breakdown of the bloodbrain barrier, acute glial activation, brain edema, and structural abnormalities of various subtypes of brain cells. These effects are also strongly enhanced when the drug is used at moderately warm environmental conditions. They implied that the mechanisms underlying environmental modulation of acute drug neurotoxicity focus on the role of brain temperature, a critical homeostatic parameter that could be affected by metabolism-enhancing drugs and environmental conditions and affect neural activity and functions.

Another important consideration is the concomitant consumption of caffeine, i.e., daily coffee intake and habitual behavior, with recreational psychostimulant drugs of abuse, may provoke varyingly severe acute adverse reactions in addition to longer-term consequences since the mechanisms by which caffeine increases the toxicity of psychostimulants include changes to body temperature regulation, cardiotoxicity, and lowering of the seizure threshold. Vanattou-Saïfoudine et al. (2012) have reviewed current notions of such caffeine-related drug interactions, placing a particular emphasis on an adverse interaction between caffeine and the substituted amphetamine MDMA which has been most recently described and characterized. The coadministration of caffeine and MDMA profoundly enhances the acute toxicity of MDMA in rats, as manifested by high core body temperature, tachycardia, and increased mortality. In addition, coadministration of caffeine enhances the long-term serotonergic neurotoxicity induced by MDMA. Their studies coincide with reports of caffeine-related drug interactions with cocaine, d-amphetamine, and ephedrine where similar mechanisms are implicated. Finally, it is an unpleasant reality that neurotoxic actions of chemical agents on humans and animals are usually studied with little consideration of the subject's nutritional status. The status of protein-calorie, vitamin, and/or mineral undernutrition levels and contingencies are linked with a range of neurodevelopmental, neurological, and psychiatric disorders, commonly with involvement of both the central and the peripheral nervous system. Undernutrition can modify risk for certain chemicalinduced neurologic diseases, and in some cases, undernutrition may be a prerequisite for neurotoxicity to surface. Spencer and Palmer (2012) have observed that combined effects of undernutrition and chemical neurotoxicity are most relevant to people in disaster areas or those with low incomes who experience, and continue to experience, chronic hunger, parasitism, and infectious disease; monotonous diets of plants with neurotoxic potential (notably cassava);

environmental pollution from rapid industrial development; chronic alcohol abuse; or prolonged treatment with certain therapeutic drugs. Finally, it is increasingly clear that drug-induced activation of central immune signaling contributes markedly to the pharmacodynamic actions of the drugs of abuse, e.g., by enhancing the engagement of classical mesolimbic dopamine reward pathways and withdrawal centers. Coller and Hutchinson (2012) have reviewed critical in vivo animal, human, biological, and molecular evidence of these central immune signaling actions of opioids, alcohol, cocaine, METH, and MDMA. These emerging notions of the central immune function signaling property of drugs of abuse present unlimited possibilities for novel therapeutic strategies and eventual interventions with opportunities to identify "at-risk" individuals through the use of immunogenetics.

# 2 Abuse Neurotoxicity at Early Developmental Stages

Brain and CNS development encompass several critical periods of vulnerability during which the developing individual is particularly sensitive to environmental neurotoxic insult, such as alcohol or nicotine (Berger et al. 1998; Rice and Barone 2000; Taranukhin et al. 2010). The effects of these drugs have been shown at both prenatal and postnatal stages of early development with regard to disruptions of both structure and function. For instance, the prenatal effects of alcohol exposure for the developing brain and CNS have been described (Goodlett et al. 2005; Riley et al. 2001; Savage et al. 2010). Purkinje cell vulnerability, for instance, occurs during postnatal days 4–6 in rodents when peak blood ethanol concentration (BEC) determines the extent of cell loss (Pierce et al. 2010). Oxidative stress is one of several mechanisms through which drugs instigate a neurotoxic action (Crews and Nixon 2009; Haorah et al. 2008). Alcohol induces lipid peroxidation in the developing brain (Kumral et al. 2005; Wang et al. 2012). Ramezani et al. (2011) observed that ethanol, administered 6 g/kg on postnatal days 4 and 5, induced lipid peroxidation, increased thiobarbituric acid reactive substance (TBARS), and decreased glutathione peroxidase levels in the cerebellum of rat pups, as well as impairing rotarod performance and locomotor activity at adolescent age of test. They observed also the neuroprotective action of 17β-estradiol, injected 30 min before ethanol, which attenuated the motor impairments and lipid peroxidation and restored antioxidant levels. Alcohol is now generally acknowledged as the leading human teratogen and, if the estimates of prevalence of fetal alcohol spectrum disorders (FASD) are correct, is one of the leading known causes of neurodevelopmental disorders in Western society (Goodlett 2010).

# 3 Alcohol Neurotoxicity

Ethanol crosses the blood-brain barrier readily and is metabolized in the brain by several enzymes in processes rendering reactive oxygen species. Excessive reactive oxygen species (ROS) that disturb the normal cellular redox state induce oxidative

stress leading to cellular damage (Hampton and Orrenius 1998). Alcohol is a neuroteratogen which induces loss of neurons in the CNS that from a neurodevelopmental aspect account for several of the symptoms expressed in fetal alcohol syndrome (Chen et al. 2008; Riley and McGee 2005). Certain regions of the developing brain are selectively vulnerable to alcohol and there are peak periods of sensitivity within these regions (Rice and Barone 2000; Stoltenburg-Didinger 1994). A single ethanol exposure in rats during the first several postnatal days (P4-5) that correspond to the human third trimester induces a huge loss of neurons in brain (Bonthius and West 1991; Hamre and West 1993); slightly later periods show greater resistance to similar exposure (Goodlett and Eilers 1997; Light et al. 2002). The window of vulnerability (also termed peak sensitivity, although the latter is more reliably linked to vision) coincides with the developmental period of synaptogenesis, also known as the brain growth-spurt period, which in rodents is a postnatal event, but in humans extends from the 6th month of gestation to several years after birth. It appears likely that the alcohol-induced disruption of neurodevelopment involves apoptotic processes implicating the Bcl-2 family of survival-regulatory proteins (Ge et al. 2004; Lee et al. 2008; Ullah et al. 2012). The proapoptotic Bax protein, within the Bcl-2 gene family, is implicated in alcohol-induced cell death in developing organisms. After neonatal ethanol administration during ages of peak sensitivity, Bax migrates to the mitochondrial membrane where it participates in initiating apoptosis (Siler-Marsiglio et al. 2005; Heaton et al. 2012a). Heaton et al. (2012b) studied the interactions between Bax and proteins of the permeability transition pore (PTP), voltage-dependent anion channel (VDAC), and adenine nucleotide translocator (ANT) of the outer and inner mitochondrial membranes. They found that after alcohol treatment, Bax proapoptotic associations with VDAC and ANT were increased, especially at peak sensitivity, and sustained at least 2 h posttreatment. Ethanol neurotoxicity in cultured neuronal preparations was abolished by pharmacological inhibition of VDAC and ANT interactions but not by a Bax channel blocker. They concluded that, following ethanol treatment, BAX initiated the apoptosis cascade through interactions with proteins of the PTP complex but not channel formation independent of PTP constituents.

Generally, chronic alcohol intake is more harmful than acute intake (Eysseric et al. 1997; Lamarche et al. 2003). The brain and CNS, and not least neurons, are major targets for the uptake and pharmacological action of alcohol, and chronic consumption of ethanol is associated with several neurodegenerative disorders (Brooks 1997, 2000; Dlugos 2009) and conditions of neurodegeneration (Qin and Crews 2012). The brain damage is characterized by cerebral and cerebellar atrophy and widespread neuronal deficits in several regions in adult chronic alcohol abusers (Bleich et al. 2003; Harper 1998; Harper and Matsumoto 2008). Alcohol-related neuropathology is associated with both excitotoxic (Krystal et al. 2003) and oxidative stress (Sakaguchi et al. 2011; Tiwari et al. 2012). Ethanol exposure produces ROS which cause oxidative stress in the brain. The excessive production of ROS during ethanol catabolism or from damaged mitochondria following ethanol exposure exacerbates the neurotoxicity, particularly during brain development. Kruman et al. (2012) have outlined mechanisms involving the accumulation of DNA

damage in the absence of cellular repair that unfolds in the expression of genomic instability and neuronal death. Genomic instability involves structural alterations that modify the standard chromosomal complement; this process may lead to a situation of aneuploidy, an abnormal number of chromosomes, through which the cells present a chromosomic number that is either higher or lower than the normal complement for the species. For example, alcohol is capable of contributing toward the initiation of cancer through the induction of genomic (chromosomal) instability and aneuploidy, which may be a result of multiple indirect mechanisms (Benassi-Evans and Fenech 2011). Alcohol interacts also with different metabolic pathways. The one-carbon metabolism (OCM) is critical for the synthesis of DNA precursors. It is also a methyl donor for methylation events. Maintenance of genomic stability is dependent upon the processes of DNA repair and methylation such that the interaction of ethanol with OCM poses the threat of DNA damage and DNA repair dysfunction (Kruman et al. 2012).

Autophagy is a carefully regulated pathway that involves the lysosomal degradation of cytoplasmic organelles or cytosolic components, a cell biological process providing a central component of the integrated stress response. Since eukaryotic cells, defined by their membrane complexity, are under constant need to adapt to changing environments (or "stressors" of varying magnitude), it is the response of such cells to these fluctuating stressors that determines whether functionality and survival or cell death occurs. Incorporated in the response to stress (assumed sublethal) is the capacity for rapid changes adapting metabolism and promoting protection. Autophagy is a major pathway mediating stress-induced metabolic adaptation and protective measures that promote species survival against neurodegenerative, neuroinflammatory, and neoplastic stressors, involving abnormal cell proliferation (Levine and Kroemer 2008; Mizushima et al. 2008). A close integration between regulatory processes in response to stress, including nutrient uptake, metabolism, cell cycle and growth control, cell fate and lineage decisions, and cellular survival and death programs, and processes regulating autophagy is maintained, thereby promoting the fundamental adaptation to environmental stressors. Impairments in autophagy lead to accumulation of damaged proteins and organelles thereby eliciting increased cellular stress. Autophagy, through the elimination of potential proapoptotic sources such as damaged mitochondria, provides a cytoprotective pathway which sets a higher threshold against global apoptosis induction (Kroemer et al. 2010). The survival function of autophagy in nutrient- or growth factor-deprived cells (Klionsky 2007) is linked to cellular repair and antipathogen defense (Melendez and Neufeld 2008). The generation of oxidative stress and mitochondrial-related changes by alcohol is an important aspect of the damaging influence of the drug (McVicker et al. 2009). Nonetheless, since autophagy exerts either a pro-survival or pro-death effect, its influence by exposure to the drug is of some importance. Von Haefen et al. (2011) have shown that ethanol downregulated the autophagy proteins, Beclin-1 and LC3-II, with enhanced apoptosis through breakdown of mitochondrial potential, upregulation of cleaved caspase-3 and PARP-1, and downregulation of antiapoptotic protein Bcl-2. They concluded that ethanol inhibited autophagy thereby enhancing vulnerability for cell death.

## 4 Nicotine Neurotoxicity

The disrupting influences of developmental neurotoxicity by drugs of abuse presents an important regulatory issue as well as a necessary area for significant public concern (Archer and Kostrzewa 2012). Despite the plethora of evidence, the developmental neurotoxicity of a huge number of compounds has yet to be determined (Judson et al. 2009). This type of neurotoxicity involves several manners of expression, including cell proliferation (Breier et al. 2008), synaptogenesis (Harrill et al. 2011), gene expression (Hill et al. 2008), neurite formation (Radio and Mundy 2008), network formation (Robinette et al. 2011), and function and regional integrity (Lai et al. 2011). Applying methods for detecting chemical-induced alterations in proliferation, apoptosis, and viability, Culbreth et al. (2012) have described viability (status of a cell to survive, grow, and multiply) changes in human neuroprogenitor cells following nicotine exposure. Despite evidence that cigarette smoking during pregnancy is associated with fetal, obstetrical, and developmental outcomes, 15-20 % of all women smoke throughout the duration of pregnancy (Andres and Day 2000; Srisukhumbowornchai et al. 2012). Maternal smoking results in myriad physical, cognitive, and behavioral effects in offspring due to prenatal exposure to nicotine. A great proportion of the adverse physiological symptoms attributed to smoking termination (craving, irritability, restlessness, anxiety, and appetite) are related to nicotine withdrawal (Glynn et al. 2009). Through its capacity to bind with cholinergic nicotinic receptors (nAChRs), nicotine dysregulates several essential biological processes such as angiogenesis, apoptosis, and cell-mediated immunity (Zeidler et al. 2007), with proapoptotic effects, in vivo and in vitro, particularly in the developing brain (Abreu-Villaca et al. 2004; Qiao et al. 2003; Roy et al. 1998, 2002). Finally, the effects of nicotine upon permeability of the blood-brain barrier may expose brain regions to potentially serious neurotoxicity (Manda et al. 2010).

Nicotine taken in by pregnant women crosses the placenta, concentrates in fetal blood and amniotic fluid, and is detectable in breast milk during lactation (Lambers and Clark 1996). The effects of nicotine replacement therapy for offspring are linked to fetotoxicity and neuroteratogenicity (Ginzel et al. 2007), as well as adverse cardiovascular, respiratory, endocrine, and metabolic outcomes (Laitinen et al. 2012; Oken et al. 2008; Syme et al. 2010). In order to model the human maternal smoking situation in the animal laboratory, a gestational nicotine exposure has been devised (Coleman 2008; Dwyer et al. 2008, 2009; Franke et al. 2007; Prins et al. 2012). Similar to prenatal smoking exposure in humans, the gestational exposure in animals dose-dependently induces low birth weight, cognitive performance deficits, and enhanced locomotor activity (Eppolito and Smith 2006). Nicotine, a major neurotoxin in cigarette smoke, crosses the placental barrier and fetal, poorly-developed, blood-brain barrier and binds to nAChRs, thereby contributing to the pathogenesis of abnormal brainstem neurochemistry. The gestational nicotine model presented also extensive neurochemical deficits that were manifested during adolescence (Franke et al. 2008; Slotkin 2008). Nevertheless, not all the prenatal nicotine alterations seem to imply permanent deficits. Santiago and Huffman (2012) have observed that although mouse mother nutrition, dam weight gain, and litter size were not significantly affected by nicotine treatment, prenatal nicotine exposure resulted in lower newborn birth weight and brain weight and length. Remarkably however, the reduction of body weight, brain weight, and brain length observed in newborn prenatal smoking exposure mice, compared to control mice, was no longer present at the 10th postnatal day. Wei et al. (2011) used a pathway-focused oligonucleotide microarray to analyze molecular mechanisms underlying how gestational exposure to nicotine alters brain development in five regions: amygdala, prefrontal cortex, nucleus accumbens, periventricular nucleus of the hypothalamus, and caudate putamen. They identified 24 cell death/survivalrelated pathways modulated significantly by gestational nicotine that were classified into three categories: (i) growth factor, (ii) death receptor, and (iii) kinase cascade, on the basis of their biological functions. Kinase cascades present a series of several protein kinases that phosphorylate and activate each other: Kinase-1 phosphorylates and activates kinase-2, in turn then phosphorylating and activating kinase 3. This cascade serves to amplify greatly a signal since each protein kinase can phosphorylate multiple substrates. Quantitative real-time PCR (polymerase chain reaction) array was applied to verify their findings through measures of 29 genes involved in cell death/survival-related pathways and indicated that gestational nicotine exposure exerted significant effects on these pathways in the brains of adolescent offspring. These effects were brain region specific and actualized through the expression of growth factors and receptors, caspases, kinases, and transcription factors. They have presented the notion that gestational nicotine exposure affects cell death and survival in adolescent offspring brains by regulating the balance between growth-factor and death-receptor pathways (Wei et al. 2011).

The apoptotic propensity of fetal and neonatal nicotine exposure and its accompanying health hazards have been described both neurodevelopmentally and pathophysiologically (Bruin et al. 2008; 2012). In this vein, it ought to be noted that, whereas there is no information on the long-term effects of developmental nicotine exposure in humans, the consensus from laboratory studies indicates that nicotine alone is the key factor underlying pathophysiology, including Type 2 diabetes, neurobehavioral defects, obesity, hypertension, impaired fertility, and respiratory dysfunction (Bruin et al. 2010). The neurotoxic agency of nicotine is not straightforward but rather complex and enigmatic. It would appear that nicotine exerts, concurrently, both positive, neuroprotective, and apoptosis-inhibiting effects (Hejmadi et al. 2003; Tizabi et al. 2005; Zhang et al. 2006) and conversely negative, neurotoxic, and apoptosis-promoting effects (Barr et al. 2007; Crowley-Weber et al. 2003; Zhao and Reece 2005) with regard to the integrity of the developing and the mature brain. Nicotine, with its specific binding to nAChR, functions as a dysregulator of essential developmental biological processes, such as cell proliferation, apoptosis, migration, invasion, angiogenesis, inflammation, and cell-mediated immunity in a wide variety of cells including fetal (regulation of development), embryonic, and adult stem cells, adult tissues, as well as invading cancer cells (Cardinale et al. 2012). In cardiomyocytes, which may offer a model cell type for examining neurotoxicity, nicotine promoted apoptosis by inducing oxidative stress and disrupting apoptosis-related gene expression (Zhou et al. 2010). The neurodevelopmental toxicity of nicotine has been described to some extent and in comparison with other real and potential neurotoxicants using zebrafish development as the laboratory model (Muth-Köhne et al. 2012). From these propensities, it seems clear that the action of the drug is multifaceted, possibly involving different neuronal circuits influencing each other through complicated interactions. Ferrea and Winterer (2009) have summarized the most important results of experiments about nicotinic neuroprotection and neurotoxicity propensities in humans and animals. In the first place, well-known modifications of cholinergic transmission during physiological (normal aging) and pathological neurodegeneration have been outlined, and in the second place, neuroprotective and neurotoxic effects of nicotine, also mentioning the underlying molecular mechanisms, have been described.

Maternal smoking exerts critically health hazards that may through indirect means induce neurotoxic actions in brain regions leading to functional deficits. Multiple pathophysiological mechanisms, e.g., respiratory control abnormalities, contribute to these defects. Smoking/nicotine during pregnancy pressurizes the maternal metabolic system and contributes to adverse pregnancy outcomes, predicting future risks for respiratory dysfunction, neurobehavioral deficits in cognitive-emotional domains, cardiovascular disease, obesity, and Type-2 diabetes (Ellis et al. 2012; Langley et al. 2012; Mamun et al. 2012; Zhu et al. 2012). "Catchup" growth is observed normally in children exposed to intrauterine smoke which was linked to subsequent childhood obesity (Chen et al. 2012). The abnormal hypothalamic gene expression of appetite regulators, e.g., downregulation of proopiomelanocortin and neuropeptide Y in the arcuate nucleus of the hypothalamus may be one expression of health hazard. Human studies indicate that maternal smoking causes changes in ventilator control, and respiratory and arousal patterns in newborn infants, decreasing ventilator drive and blunted hypoxic ventilator response (Ueda et al. 1999). There exists a dose-dependent increase in the frequency and duration of obstructive respiratory events in term infants born to mothers who smoked (Kahn et al. 1994). Prenatal exposure to cigarette smoke is associated with decreases in arousal responses after obstructive events (Tirosh et al. 1996). Preterm infants born to smoking mothers showed increased apneic events and decreased respiratory arousal (Sawnani et al. 2004). Nicotine, despite the presence of other chemicals in cigarette smoking, provides the major factor underlying alterations in respiratory control. Nicotine-exposed neonatal rats showed lower minute ventilation on room air thereby evidencing nicotine effects on eupneic, normal, good breathing, ventilation (St. John and Leiter 1999), showing lower tidal volume and higher respiratory frequency (Hafstrom et al. 2002a). Prenatal nicotine attenuates the hypoxic ventilator response (Bamford and Carroll 1999; Hafstrom et al. 2002b; Milerad et al. 1995). Prenatal nicotine exposure is accompanied by a decrease in the magnitude of hypoxic ventilator depression, the component of hypoxic ventilator response that activates the PDGF-B receptor (platelet-derived growth factor- $\beta$  receptor) and its downstream antiapoptotic cascade in the caudal brainstem of developing rats (Gozal et al. 2000).

Simakajornboon et al. (2010) examined the effect of prenatal nicotine exposure on PDGF- $\beta$  receptor activation and the subsequent activation of downstream antiapoptotic processes through the Akt/BAD pathway (Akt promotes cell survival through phosphorylation of BAD) which in itself is inhibited by stress (Zundel and Giacco 1998). They concluded that prenatal nicotine exposure attenuated the phosphorylation of PDGF- $\beta$  receptor, Akt, and BAD-136 during hypoxia in the caudal brainstem of developing rats, suggesting the elevated apoptosis in this region and increased vulnerability of neural cells in the respiratory control area.

#### 5 Alcohol-Nicotine Combinations

Adolescence, characterized as a period of vulnerability (Grant 1998), is a critical period with likely debut for both smoking and alcohol (National Institute on Drug Abuse 1998; Spear 2000), and adolescent "drinkers" are more likely to be "smokers" than "non-drinkers," and vice versa (Chen and Kandel 1995; Van Zundert et al. 2012; Zhang et al. 2012); the consequences are much more serious with pregnant mothers (McDonnell-Naughton et al. 2012). Nicotine, by itself, is damaging to the adolescent brain (Abreu-Villaca et al. 2003). Adolescents appear to be consuming larger quantities of cigarettes and alcohol earlier in life (Perkonnig et al. 2006) with a strong relationship between onset of tobacco use at an early age and alcohol addiction (Burns and Proctor 2012; Muñoz et al. 2012). Smoking prevalence among individuals under treatment for alcoholism is 3-4-fold that of the general population (Batel et al. 1995). It has been shown that nicotine and ethanol interact during adolescence among mice to affect cognitive performance (Abreu-Villaca et al. 2007) and affective status (Abreu-Villaca et al. 2008), during both exposure and withdrawal. Hippocampal cell death and reduced densities of neuronal and glial cells were observed during adolescent nicotine or ethanol exposure whereas the drug combination caused less significant effects (Oliveira-da-Silva et al. 2009). The effects of nicotine and/or ethanol upon several regions of the hippocampus were investigated (Oliveira-da-Silva et al. 2010); these included the granular layer of the dentate gyrus, molecular layer, CA1, CA2, and CA3 of C57/B16 mice, using TUNEL assay. There were increases in the numbers of TUNEL + cells in both the nicotine and ethanol groups, with less severe effects in the nicotine-ethanol combination group. Their results point toward the deleterious effects of nicotine and/or ethanol that were reversed during prolonged withdrawal.

Both animal and human studies have described tolerance, consumption, relapse, and behavioral interactions between ethanol and nicotine administration. Nevertheless, the interactions of these drugs with particular reference to developmental and adolescent neurotoxicity, ethanol withdrawal in adulthood for subjects who have an adolescent history of using these drugs, and their combined effects on neuroimmune signaling and mobilization remain neglected although attention emerges (Durazzo et al. 2007). For example, Riley et al. (2012) have investigated nicotine's influence on ethanol withdrawal seizures in two different age groups of male C3H mice. Adolescent and adult male C3H mice, beginning at postnatal day 28 or 70, respectively, were subjected to a 7-day chronic exposure to ethanol only, ethanol plus nicotine, nicotine only, or vehicle treatment. Six weeks later, all the groups were subjected to chronic exposure to ethanol vapors and the severity of their ethanol withdrawal seizures was assessed by handling-induced convulsions. An adolescent exposure to chronic nicotine resulted in an exacerbation of ethanol withdrawal seizures in adulthood. Given this, adolescence may contain a neurophysiological critical period that is sensitive to nicotine and which may result in an altered response to ethanol dependency in adulthood. These findings have serious implications for the long-term consequences following co-abuse of these drugs during adolescence. The synergistically detrimental effects of combined alcohol and nicotine exposure upon neurodevelopment

and brain and CNS integrity ought not to be underestimated (Ramlochansingh

et al. 2011). The effects of nicotine on neural mechanisms modulating respiratory and cardiovascular functioning outlined above become rather more noteworthy with the neurophysiological interactions that arise in combination with alcohol. The prenatal exposure to high levels of nicotine or alcohol separately, e.g., in tribal groups of the North American Plains Indians, has been considered to constitute a major factor contributing to sudden infant death (Gunn et al. 2000; Isayu et al. 2002). In an autopsy study of the Aberden Area Infant Mortality Study, Duncan et al. (2010) investigated the effects of cigarette smoke and alcohol upon the distribution and binding of nAChRs in the brainstem of sudden death cases (SIDs). The expression of nAChRs in the brain is widespread, primarily presynaptic. These receptors, contributing to neurotransmitter release, neurite outgrowth and differentiation, and neuronal survival (Gotti and Clementi 2004; Torrao and Britto 2002), are ligand-gated ion channels, pentameric in structure, and comprised of homomeric or heteromeric combinations of 12 genes [ $\alpha$ 2-10,  $\beta$ 2-4] (Landgren et al. 2011). The <sup>3</sup>H-nicotine radioligand was chosen since it binds, with varying affinity, to all subunits of nAChRs but preferentially to the  $\alpha_4\beta_2$  subunit to which nicotine in cigarette smoke binds (Benwell et al. 1988). <sup>3</sup>H-nicotine binding in five nuclei related to cardiorespiratory control decreased with increasing number of drinks (Duncan et al. 2008). In three mesopontine nuclei critical for arousal, there were reductions in binding for controls exposed to cigarette smoke compared with controls not exposed (ibid). The authors indicate that maternal smoking and alcohol affected <sup>3</sup>H-nicotine binding in the infant brainstem, irrespective of cause of death. Durazzo et al. (2004) used human in vivo proton magnetic resonance spectroscopic imaging to observe that chronic cigarette smoking exacerbated chronic alcohol-induced neuronal injury and cell membrane damage in the frontal lobes of recovering alcoholics (14 smokers and 10 nonsmokers) in treatment compared with 26 light-drinking controls (7 smokers and 19 nonsmokers). They found independent adverse effects on neuronal viability and cell membranes in the midbrain and on cell membranes of the cerebellar vermis, with higher smoking levels being associated with metabolite concentrations in the selected subcortical structures.

## 6 Conclusion

Substance abuse, as above with ethanol, nicotine, or both, is associated with levels of neurotoxicity encompassing most systems of the brain and CNS. The transition from occasional-recreational use to chronic abuse and addiction presents serious psychiatric obstacles that allow few avenues for effective and definitive treatment since most individuals relapse, even after long periods of abstinence. The elucidation of mechanisms through which these substances exert neurotoxicity and mediate addiction remains an essential requirement and mostly involves altered gene function (Kovatsi et al. 2011). Thus, the increasing evidence derived from epigenetics implicating the epigenome in situations of altered gene function may guide through to an understanding of etiopathogenesis. Nevertheless, despite the identification of several genes linked to alcoholism, the genetic basis remains a mystery; advances pursued through the development of transcriptomics, epigenomics, and proteomics may eventually explain genetic susceptibility to alcohol (Buscemi and Turchi 2011). It is expected that the eventual understanding of fetal alcohol syndrome disorders requires (1) the identification of genetic and epigenetic modifications that may be predictive of the neurobehavioral and neurobiological dysfunctions in offspring induced by gestational alcohol exposure and (2) the determination of the relationship structural brain alterations induced by gestational alcohol exposure and the functional-epigenetic outcomes in the offspring (Reynolds et al. 2011). Both the major problem of adverse drug reactions and related neurotoxicity (Kacevska et al. 2011) and the influence of chronic nicotine administration prior to cocaine abuse that exacerbates the epigenetic, gene expression, and electrophysiological and neurobehavioral effects during the transition from acute to chronic responses to cocaine (Volkow 2011) may be linked to the addictive process (Archer et al. 2011, 2012; Archer and Kostrzewa 2013; Blum et al. 2011).

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# **Experimental Approach to Alzheimer Disease**

M. Salkovic-Petrisic, S. Hoyer, and Peter Riederer

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### Abstract

This review on current experimental models of Alzheimer disease is based on human postmortem findings showing keystone markers for a pathology within the  $\beta$ -amyloid transduction cascade as well as pathology in the mechanism of phosphorylation of  $\tau$ au protein. Evidence for risk factors triggering this devastating disease focuses on type II diabetes. Therefore, modelling Alzheimer disease tries to get profound knowledge of these underlying mechanisms by studying experimental animal models. Here will be discussed in detail two

M. Salkovic-Petrisic

S. Hoyer

P. Riederer (🖂)

Department of Pharmacology and Croatian Institute for Brain Research, School of Medicine, University of Zagreb, Zagreb, Croatia

Department of Pathology, University of Heidelberg, Heidelberg, Germany

University of Würzburg, Department of Psychiatry, Psychosomatics and Psychotherapy, and National Parkinson Foundation Centers of Excellence for Neurodegenerative Diseases Research, Laboratory for Clinical Neurochemistry, Würzburg, Germany e-mail: peter.riederer@mail.uni-wuerzburg.de

pharmacological models, the one mirroring type II diabetes-induced AD pathology by streptozotocin and its influences on the insulin/insulin receptor cascade as well as  $\beta$ -amyloid and  $\tau$ au pathologies. Behavioral studies give evidence that this model is currently the most detailed described AD model. While transgenic mouse models, like the APP Tg2576 model, demonstrate  $\beta$ -amyloid plaque formation and impaired memory rather in old age, streptozotocin is able to aggravate the process of pathology so that AD pathology is seen months earlier. This indicates a profound interaction of AD pathology with the insulin/insulin receptor cascade and pathobiochemistry.

Modelling cholinergic deficits has been done by using the cholinotoxin AF64A. This model reflects changes in the acetylcholine metabolism as well as in monoaminergic transmitter systems, and behavioral studies are in line with AD-based impairment of learning and memory. However, ß-amyloid and tau pathologies as well as insulin-resistant brain state have not been studied in detail.

#### **Keywords**

AF64A model • Alzheimer disease • Dementia model • Streptozotocin • Transgenic dementia models • Type II diabetes

## 1 Introduction

Any model of a disease should reflect the core abnormalities of that disease in morphobiological and clinical terms. In case of Alzheimer disease (AD), it has to be taken into account that AD is not a single disorder from a nosological point of view. A very small proportion of 513 families worldwide (by August 2011) are suffering from AD caused by missense mutations: in the amyloid precursor protein (APP) gene on chromosome 21 (89 families ~ 17.3 %), in the presenilin gene 1 on chromosome 14 (402 families ~ 78.4 %), and in the presenilin gene 2 on chromosome 1 (22 families ~ 4.3 %) (http://www.molgen.ua.ac.be/admutations). These currently known three mutations are involved in both the overexpression and the abnormal cleavage of APP and the increased formation of the APP derivative  $\beta$ -amyloid (A $\beta$ 1–40, A $\beta$ 1–42/43), a keystone for the amyloid hypothesis of AD (Sandbrink et al. 1996; Hardy and Selkoe 2002; Selkoe 2003). Clinically, the dementia symptoms may become evident as early as in the third decade of life.

In contrast, in sporadic AD (sAD), which approximately more than 15 million people worldwide suffer from (Blennow et al. 2006), no such mutations were found as yet, i.e., the mutation-related stimulus is lacking, and its role in the causation of sAD remains enigmatic (Hoyer 1993; Joseph et al. 2001). Also, in longitudinal studies on aged nondemented, cognitively normal subjects, abundant neuritic plaques and neurofibrillary tangles were found in the brain (Crystal et al. 1988; Davis et al. 1999; Snowdon 1997). Moreover, the decline of the mental status starting in later life and followed over several years was demonstrated not to be

accompanied by changes in the numerical density of plaques and tangles (Bennett et al. 1993; Davies et al. 1987; Mann et al. 1985). Finally, it was the neuronal loss rather than neuritic plaques and neurofibrillary tangles that directly contributed to cognitive damage (Gomez-Isla et al. 1997). Also, in postmortem sAD brain, AB42 immunization over 84 weeks induced a marked reduction of amyloid plaques. However, this clearance did not prevent both the progression of dementia and severe end-stage dementia before death (Holmes et al. 2008). Overall, the above findings clearly indicate that sAD is due to risk factors other than mutations in the APP gene or other genes.

Among the risk factors under discussion are, first, age as a leading one (Evans et al. 1989; Ott et al. 1995); second, a single nucleotide polymorphism found in the gene coding for 11*B*-hydroxysteroid dehydrogenase I, which is associated with a sixfold increased risk for sAD (de Quervain et al. 2004); and, third, allelic abnormalities of the APO-E 4 gene on chromosome 19 (Mahley and Rall 2000; Tanzi and Bertram 2001). People carrying one copy of this allelic abnormality (around 24 % of the population) or carrying two copies (around 2–3 % of the population) were found to show drastic abnormalities in the cerebral metabolic rate of glucose (Corder et al. 1993), i.e., the nutritional state of the brain may be severely perturbed under these conditions.

The latter data may indicate that the cerebral glucose metabolism may be of pivotal significance. In the normal mature mammalian brain, oxidative glucose/ energy metabolism is largely under the control of the neuronal insulin signal transduction cascade (Hoyer 2004a; Maurer and Hoyer 2006). This metabolic pathway is the source of acetyl-CoA necessary for the formation of the neurotransmitter acetylcholine and of ATP which drives nearly all cellular and molecular processes (for a review, see Salkovic-Petrisic et al. 2009). In patients suffering from very early sAD, the cerebral glucose metabolism was found to be reduced in the posterior cingulate cortex (Minoshima et al. 1997) whereby the impairment of cerebral glucose metabolism was found to be of predictive significance (Mosconi 2005). As sAD progresses, the abnormalities in cerebral glucose utilization grow more severe, whereas cerebral blood flow and oxygen consumption of the brain were found to be much less perturbed (Foster et al. 1984; Hoyer et al. 1991; Mielke et al. 1994), i.e., the abnormal glucose metabolism in the brain may be in the center of metabolic abnormalities in sAD, although the glucose concentration in arterial blood was found to be unchanged (Fig. 1).

### 2 Relevant Animal Models

Based on the latter, cerebral glucose metabolism may be experimentally damaged via the insulin receptor in the brain by two different ways: first, by chronic elevation of the circulating corticosterone (cort) concentration and, second, by direct functional damage by the diabetogenic substance streptozotocin (STZ) applied to the brain only.

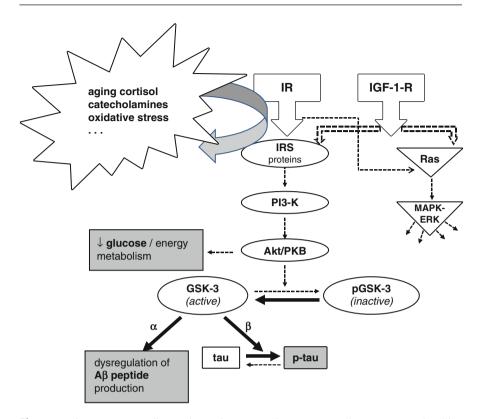


Fig. 1 Insulin receptor signalling in the insulin-resistant brain state. Insulin receptor (IR) signalling in the brain may be affected by various factors: physiological and pathophysiological conditions like aging, cortisol, catecholamines, and oxidative stress (as reviewed elsewhere, Hover 2004, Salkovic-Petrisic and Hoyer 2007). They may act at the level of the IR and its tyrosine kinase activity and/or at any other element downstream the IR signalling cascade (insulin receptor substrate/IRS) leading to its dysfunction which is finally expressed as the insulin-resistant brain state (IRBS), found in both sAD humans and respective animal models (Hoyer 2004; Salkovic-Petrisic and Hoyer 2007). Namely, the target elements downstream the IR signalling pathways, in particular the PI3K pathway parameters presented at the figure, behave like they are resistant to the influence of insulin due to the reduction/ loss of physiological insulin functioning. Low activity of AKT/PKB protein kinase, induced by upstream IR signalling malfunctioning, leads to (i) a decrease in glucose/energy metabolism via glucose transporters and (ii) less phosphorylation and by that activation of glycogen synthase kinase-3 (GSK-3) isoforms  $\alpha$  and  $\beta$ . Activated GSK-3 $\alpha$  is involved in the dysregulation of  $\beta$ -amyloid peptide (A $\beta$ ) production, while activated GSK-3 $\beta$  favors hyperphosphorylation of tau protein, all being a hallmark of sAD pathology. In the STZ-icv rat model, IRBS is induced by the icv administration of low doses of STZ, a drug selectively toxic to glucose transporter-2 and IR expressing cells (Szkudelsky 2001). Solid arrows, function stimulated; dashed arrows, function inhibited

### 2.1 Corticosterone Model

It has been well documented that chronic elevation of cort in circulating blood induced a strong increase of cort in both cerebral cortex and hippocampus associated with both learning and memory deficits (Wüppen et al. 2010;

Hoyer and Lannert 2008; Feldmann et al. 2008). This state can be achieved by prenatal stress of (rat) dams during the last week of pregnancy and postnatal daily handling of the pups during the first 3 weeks of life. However, cognitive alterations in Y maze and radial maze were observed only in later life (Steen et al. 2005). Chronic glucocorticoid administration (as a stress model) reduced insulin-degrading enzyme (IDE) protein levels and IDE mRNA in different regions of aged macaque brain associated with increased AB42 levels relative to AB40 levels (Rivera et al. 2005).

Exposure of 3 T3-L1 cells to dexamethasone resulted in decreased insulin stimulation of glucose uptake (Moloney et al. 2010). This generation of an insulin-resistant brain state was found to be characterized by reduced gene expression of insulin and insulin receptor density and associated by increased tau protein gene expression (Vallee et al. 1999). Additionally, cerebral glucose metabolism was shown to be markedly reduced by corticosterone (Hoyer and Lannert 2008; Kulstad et al. 2005).

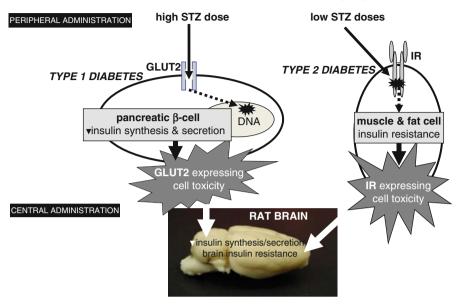
As related to human beings, a disinhibition of the HPA axis causes hypercortisolemia in sAD (Grünfeld et al. 1981), and hypercortisolemia may be supplemented by 11ß-hydroxysteroid dehydrogenase to amplify glucocorticoid action in sporadic AD (Grünfeld et al. 1981) to increase its risk.

### 2.2 Streptozotocin Model

With respect to insulin/insulin receptor (IR) in sAD brain, a decrease was found in both insulin concentration and tyrosine kinase activity along with an increase in IR density (Frölich et al. 1998). More recent studies demonstrated a drastic reduction of both the expression of insulin and IGF-1 and the respective receptors in different areas of the AD brain accompanied by a reduced insulin signal transduction capacity (Plaschke et al. 1996), whereby the expression of insulin, IGF-1 and IGF-2, and their receptors decreased with advancing AD (Pascualy et al. 2000). Intraneuronally, alterations were found in a predominance at internal and nuclear sites and in a diminution at cytoplasmic and dendritic sites (Frölich et al. 1998). These above findings may point to the neuronal insulin/insulin receptor system as the site of the predominant and early abnormality in sAD, i.e., an insulin-resistant brain state may be assumed.

It is difficult to establish an experimental animal model that would faithfully mimic the developmental pathology of AD in humans, in particular the pathology of the prevailing sporadic form, the causation of which is still unclear. The streptozotocin (STZ) model that will be described here has actually been reported in literature more than 20 years ago (Mayer et al. 1990), but the proposal that it could be considered as a model for the sAD came some 8 years afterwards (Lannert and Hoyer 1998).

STZ (2-deoxy-2-(3-(methyl-3-nitrosoureido)-D-glucopyranose)) is a betacytotoxic drug which, following peripheral (parenteral) administration at high doses, selectively destroys insulin-producing/insulin-secreting  $\beta$  cells in the pancreas and causes type I diabetes mellitus in adult animals (Szkudelski 2001). Treatment with



**STREPTOZOTOCIN** 

2-deoxy-2-({[methyl(nitroso)amino]carbonyl}amino)-β-D-glucopyranose

**Fig. 2** *Mechanism of action of streptozotocin.* Due to its chemical structure (2-deoxy-2-(3-(methyl-3-nitrosoureido)-D-glucopyranose)), streptozotocin (STZ) is a selective substrate of the glucose transporter-2 (GLUT2) which at the periphery enables entry of this compound into the insulin-producing/insulin-secreting cells, the highest expression of which is on the membrane of pancreatic  $\beta$  cells (Szkudelsky 2001). Following peripheral (parenteral) administration at high doses, STZ selectively destroys insulin-producing/insulin-secreting  $\beta$  cells in the pancreas and causes type I diabetes mellitus in adult animals (Szkudelsky 2001). Treatment with low to moderate doses of STZ causes insulin resistance (Blondel and Portha 1989) via a decrease in autophosphorylation (Kadowaki et al. 1984) and an increase in total number of insulin receptor (IR), but with little change in phosphorylated IR- $\beta$  subunit (Giorgino et al. 1992). The mechanism of central STZ action and its target cells/molecules in the brain have not been identified yet, but given the presence and the regionally specific distribution of GLUT2 and IR in the mammalian brain, similar mechanism may have been speculated to induce the insulin-resistant brain state following the STZ-icv administration

low to moderate doses of STZ has been demonstrated to cause insulin resistance (Blondel and Portha 1989) via a decrease in autophosphorylation (Kadowaki et al. 1984) and an increase in the total number of IRs, but with little change in phosphorylated IR- $\beta$  subunit (Giorgino et al. 1992) (Fig. 2). Considering this selectivity to insulin system-related cells and a presence of insulin and IRs in the brain, the idea of the first experiments performed in the 1990s was that the central application of low STZ doses (up to 100x lower than peripheral ones) into the lateral ventricles (icv) might be associated with a kind of local, cerebral diabetes which might be reflected on behavior and brain neurochemistry. Among all changes, cognitive deficits were demonstrated first in STZ-icv-treated

rats (Mayer et al. 1990) and were followed by reports of changes in brain glucose metabolism (Nitsch and Hoyer 1991; Duelli et al. 1994). In spite of that, it has been repeatedly demonstrated that central STZ administration causes neither systemic metabolic changes nor diabetes mellitus (as reviewed by Salkovic-Petrisic and Hover 2007). Generally, in the research published so far, STZ has been administrated icv into one or both ventricles of adult rats, mostly in a 1–3 mg/kg dose. There are only a few reports of intracortical administration of low STZ doses  $(40 \ \mu g/kg)$  to 3-day-old rat pups (Lester-Coll et al. 2006; de la Monte et al. 2006) and recent reports of STZ-icv administration to mice in doses similar to those in rats (Plaschke et al. 2010a; Kumar et al. 2010; Pinton et al. 2011). Variation in the susceptibility of individual animals as a characteristic feature of STZ-icv treatment has been reported by some authors (Blokland and Jolles 1993, 1994; Prickaerts et al. 2000). STZ-icv-induced effects have not been related to the direct nonspecific toxic effect of STZ at the site of drug administration because identical biochemical changes were found in the left and right striatum after unilateral administration of STZ to the right lateral cerebral ventricle (Salkovic-Petrisic et al. 1995). Further support to the specific effect of STZ within the brain has come from a number of experiments which have shown that changes in neurochemistry and morphology are not generalized all over the brain but regionally specific and pronounced at most in the hippocampus (as reviewed by Salkovic-Petrisic and Hoyer 2007).

Due to its chemical structure, STZ is a selective substrate of the glucose transporter GLUT2 which at the periphery enables entry of this compound into the insulin-producing/insulin-secreting cells, the highest expression of which is on the membrane of pancreatic  $\beta$  cells (Szkudelsky 2001). The mechanism of central STZ action and its target cells/molecules in the brain have not been identified yet, but given the presence and the regionally specific distribution of GLUT2 in the mammalian brain as well as its expression in glucose-sensing brain neurons (Brant et al. 1993; Leloup et al. 1994; Ngarmukos et al. 2001; Arluison et al. 2004a, b), similarity to the peripheral mechanism and/or cellular targets seems likely. STZ is a cytotoxic compound, and in vitro experiments on  $\beta$  cells show that within the cell it causes alkylation of  $\beta$ -cell DNA which triggers activation of poly ADPribosylation, leading to depletion of cellular NAD<sup>+</sup> and ATP (Szkudelski 2001). Decreased levels of ATP have indeed been reported in the particular brain regions following STZ-icv treatment (Nitsch and Hoyer 1991; Lannert and Hoyer 1998), further supporting the similarity between the peripheral and the central STZ mechanisms of action.

The 20-year-long research of STZ-icv rat model has extensively documented several behavioral, neurochemical, and structural features which resemble those found in human sAD patient brain post-mortem. However, knowledge about glucose metabolism and cholinergic deficit induced by STZ-icv administration and behavioral changes has not made much progress over the years. Nevertheless a significant contribution to the characterization of this model has been provided by revealing of the alterations in the brain insulin system and tau and  $\beta$ -amyloid (A $\beta$ ) pathologies, which will be elaborated further on in this text.

The main clinical feature in sAD patients is progressive memory loss, and the representative experimental model of this disease should manifest similar cognitive changes. Deficits in learning and memory functions, usually recorded in the Morris water maze swimming and passive avoidance tests, have been consistently reported and observed as early as 2 weeks after the STZ-icv administration, persisting up to 12 weeks posttreatment (the longest observational period reported until now) (Lannert and Hoyer 1998; Salkovic-Petrisic et al. 2006; Shoham et al. 2007; Grünblatt et al. 2007). The former test is measuring the spatial learning and memory function, while the latter is testing the memory acquisition of the painful stimulus which should be avoided. By means of these tests, cognitive deficits have been found regardless of age in both 1-2-year- and 3-month-old rats (as reviewed by Salkovic-Petrisic and Hover 2007). Some correlation between the spatial discrimination performance in the Morris water maze task and the decrease in hippocampal choline acetyltransferase (ChAT) activity has been found in STZ-icy-treated rats (Blokland and Jolles 1993, 1994), although no signs of reduction in specific cholinergic markers were detected in the cortex or hippocampus prior to cognitive deficit appearance (Shoham et al. 2007). The exact mechanism by which STZ-icv treatment impairs cognitive functions has still not been elucidated, but factors like direct neurotoxicity, brain glucose/energy and cholinergic deficits, oxidative stress, and insulin-resistant brain state, all found in the STZ-icv rat model, may form the biological basis for the marked reduction in learning and memory capacities found in this model.

At the neurotransmission level, similarity between the human sAD and the STZ-icv rat model has been demonstrated in altered cholinergic neurotransmission in the brain. Significant increase in the activity of acetylcholinesterase (AChE), the enzyme responsible for a rapid acetylcholine degradation, has been found from 1 to 3 weeks following the STZ-icv treatment in adult rats (Sonkusare et al. 2005; Ishrat et al. 2006; Tota et al. 2011), rat pups (Lester-Coll et al. 2006; de la Monte et al. 2006), and mice (Kumar et al. 2010), generally associated with a decrease in the activity of ChAT, the enzyme involved in acetylcholine synthesis (Hellweg et al. 1992; Blokland and Jolles 1993, 1994; Prickaerts et al. 1999; Terwel et al. 1995).

Glucose represents the primary energy source for the brain, and brain neurons are unable to synthesize or store glucose but depend on glucose transport across the blood-brain barrier and then into the neurons. Three weeks following the STZ-icv administration, concentrations of glucose and ADP, as well as glycogen levels, were increased in the cerebral cortex (Nitsch and Hoyer 1991), and glucose metabolism was significantly decreased (44 %) (Pathan et al. 2006). Six weeks following STZ-icv treatment, reduced glucose utilization (up to 30 %) was found in 17 of 35 brain areas, particularly the frontal, parietal, sensory motor, auditory, and entorhinal cortices and in all hippocampal subfields (Duelli et al. 1994). Additionally, significant decreases in activities of glycolytic key enzymes were found in the brain cortex and hippocampus (Plaschke and Hoyer 1993). Recent data has demonstrated in vivo spatial distribution of glucose hypometabolism induced by STZ-icv administration (2 mg/kg) in cynomolgus monkeys similar to that at early

stages of sAD patients, which in the monkeys persists both 6 and 12 weeks following the STZ-icv treatment (Heo et al. 2011).

STZ generates intracellular free radicals: nitric oxide (NO) and hydrogen peroxide (Szkudelski 2001). Estimations of oxidative stress induced by STZ-icv treatment commonly utilize the measurement of malondialdehyde levels (MDA), a product of lipid peroxidation used as an indicator of free radical generation, and glutathione levels, an endogenous antioxidant that scavenges free radicals and protects against oxidative stress. Significant elevations of MDA levels and decreased glutathione levels have been found in the brain up to 8 weeks following the STZ-icv treatment (Sharma and Gupta 2001, 2002; Ishrat et al. 2006; Pathan et al. 2006; Shoham et al. 2007; Kumar et al. 2010; Saxena et al. 2011) demonstrating resemblance to the oxidative stress condition in the human sAD patients post-mortem (Götz et al. 1992; Gsell et al. 1995).

Morphological changes in the brain of the STZ-icv rat model have been investigated in much lesser extent than the behavioral and neurochemical ones. The earliest study published on that suggested that a single STZ-icv dose (1.5 mg/kg) might cause neuronal damage indicated by a reduction of 30–50 % in the weight of the septum (Terwel et al. 1995). The first more direct evidence of specific neuronal damage after the STZ-icv injection has been provided by the group of Weinstock and co-workers. Using silver impregnation staining, they found that three consecutive STZ-icv injections (in total 3 mg/kg dose) caused degeneration in the fornix, a loss of white matter, and occasional clusters of myelinated axons with distorted morphology, 40 days after the first injection (Weinstock and Shoham 2004). At that posttreatment time point, selective injury to myelin and axons in the fornix and hippocampus was found in association with the activation of microglia, as demonstrated by the increased expression of the glial fibrillary acidic protein (GFAP), a marker of astrogliosis (Prieckaerts et al. 2000). The 3rd ventricle was found enlarged by 100-150 % because of a loss of ependymal cells and damage to hypothalamic periventricular myelin (Shoham et al. 2003). Their work published afterwards indicated that activated microglia and astrocytes found in the cortex, around the cannula penetration area, in the hippocampal CA1 region, corpus callosum, and medial and lateral septum, could be found already 1 week after a single STZ-icv injection (1.5 mg/kg), while 8 weeks after the injection, the whole area of the cannula penetration site contained scar tissue that was filled with tightly packed nonreactive astrocytes and no reactive microglia around (Shoham et al. 2007). Inflammatory processes, reduced neuronal density, and axonal neurotoxicity following the STZ-icv treatment have also been confirmed by recent reports (Dhull et al. 2012; Diwu et al. 2011) in addition to the report of decreased hippocampal neurogenesis found 1 month following the treatment (Qu et al. 2012). Reduced expression of neuronal- and oligodendroglia-specific genes and increased expression of genes encoding GFAP and microglia-specific proteins were also found in rat pups administered with STZ intracortically (STZ-ic) (Lester-Coll et al. 2006).

With regard to the ultrastructural changes induced by the STZ-icv treatment, a significant enlargement of the Golgi apparatus in the frontoparietal cortical neuronal cells caused by 2x-fold expansion of the trans-Golgi segment of the cellular protein secretory pathway was found 3 weeks after STZ-icv injection (Grieb et al. 2004). Considering that proamyloidogenic processing of  $\beta$ -amyloid precursor protein may occur preferentially in the trans-Golgi segment, the observed early response of neuronal ultrastructure might be of importance for predisposing cells to form  $\beta$ -amyloid deposits.

Demonstration of structural changes in the brain of the STZ-icv-treated rats manifested as AB pathology has been a great step in the process of the characterization and validation of this non-transgenic sAD model. Pathological Aß aggregation in the form of cerebral amyloid angiopathy was visualized by Congo red staining in meningeal capillaries but not earlier than 3 months after the STZ-icv injection (Salkovic-Petrisic et al. 2006) and was further supported by positive  $A\beta_1-42$  immunolabelling found in the blood vessel wall at that time point (Salkovic-Petrisic et al. 2011). In the recently published paper, thioflavine-S and Congo red staining revealed diffuse congophilic deposits in the wall of meningeal and cortical blood vessels both 6 and 9 months after the STZ-icv treatment (Salkovic-Petrisic et al. 2011), suggesting a process of progression of the cerebral amyloid angiopathy. However, no diffuse or compact amyloid plaques and neurofibrillary tangles could be detected in the rat brain up to 3 months after icv STZ treatment (Salkovic-Petrisic et al. 2006), while some preliminary work of the same authors suggests a possibility of development of the primitive plaque-like formations appearing from 6 month post-STZ-icv treatment onward (unpublished data). A positive AB42 immunosignal has been found in the brain of the STZ-intracerebrocortically treated rat pups (Lester-Coll et al. 2006). These results provide further evidence of the resemblance of the STZ-icv rat model to the human sAD patient pathophysiology and strongly suggest that there is a need to characterize the post-STZ-icv time dependency of the development of amyloid pathology in the brain of this non-transgenic model in which clear differences in comparison to the APP transgenic mice models become obvious.

None of the previously explored alterations were able to provide a missing link to the primary event which could initiate the pathological hallmarks of sAD, hyperphosphorylated tau protein in neurofibrillary tangles, and aggregated  $A\beta$ peptide in amyloid plaques. Evidence has gathered indicating that central STZ administration induces region-specific and, for some parameters, time-progressive dysfunction in the rat brain insulin signalling, i.e., an insulin-resistant brain state (IRBS) which highly resembles those found in sAD patients post-mortem (Hoyer 2004; Cole and Frautschy 2007; Qiu and Folstein 2006; Watson and Craft 2006; Wada et al. 2005), suggesting the IRBS might be the missing link. Insulin in the brain binds to IRs which have been found functionally linked to improved cognition, in particular general and spatial memory (Zhao et al. 1999; for review Park 2001; Zhao et al. 2004). Activated IR tyrosine kinase, in addition to the protein kinase (MAPK) mitogen pathway, activates also the phosphatidylinositol-3 kinase (PI3K) pathway and its downstream elements (Johnston et al. 2003), protein kinase B (Akt/PKB) which phosphorylates and consequently inactivates both  $\alpha$  and  $\beta$ cytosolic forms of glycogen synthase kinase-3 (GSK-3) (Cross et al. 1995), involved in regulation of APP metabolism and AB peptides production (Phiel et al. 2003) and in tau protein phosphorylation (Ishiguro et al. 1993), respectively. IRBS in the STZ-icv rat model is manifested as decreased expression of the insulin genes 1 and 2, as well as the IR gene mostly pronounced in hippocampus but present also in the cerebral cortex, 2 and 12 weeks following the drug administration (Agrawal et al. 2011; Grünblatt et al. 2007). This is followed by decrement in IR- $\beta$  protein level and unchanged Akt/PKB protein expression at 4 weeks but decreased by 12 weeks posttreatment (Salkovic-Petrisic et al. 2006). Recent data demonstrates significant decrease in IR protein expression, phosphorylation of its downstream substrate (IRS-1), and Akt/PKB in hippocampal CA3 region of STZ-icv-treated rats already 2 weeks following the treatment compared to the control rats subjected to cognitive training (Agrawal et al. 2011). Decreased phosphorylation of PI3K has been also demonstrated in cerebral cortex 3 weeks following the STZ-icv treatment in rats (Deng et al. 2009). Downstream the IR–PI3K pathway, IRBS is manifested by unchanged total GSK-3 $\beta$  levels and decreased p-GSK-3 $\beta$  levels with consequently decreased p-GSK-3/GSK-3 ratio in the hippocampus and cerebral cortex, found 2, 3, and 12 weeks following STZ-icv treatments, indirectly suggesting increased GSK-3ß activity (Salkovic-Petrisic et al. 2006; Deng et al. 2009; Ponce-Lopez et al. 2011). In line with dysfunction in the upper part of IR–PI3K signalling pathway manifested already 2 weeks after the STZ-icv treatment, IRBS seems to eventually progress to its lower parts with increased hippocampal expression of tau protein and the p-tau/tau ratio being manifested 4 and 12 weeks following STZ-icv treatment (Salkovic-Petrisic et al. 2006; Grünblatt et al. 2007). Similar IRBS condition with altered IR–PI3K signalling pathway has been also found in STZ-intracortically treated rat pups (Lester-Coll et al. 2006). Recent data demonstrated that the STZ-icv treatment effects on the IR–MAPK pathway might be also time and region dependent as decreased levels of phospho-ERK were found in the cerebrum 3 weeks after the treatment (Deng et al. 2009) while phospho-ERK and its upstream element, phospho-Shc, were found unchanged 2 weeks after the treatment in the hippocampus (Agrawal et al. 2011). Disturbances in insulin action, IR function, and downstream signalling pathways have been found post-mortem in the brain of sAD patients (Hoyer 2004; Cole and Frautschy 2007; Qiu and Folstein 2006; Watson and Craft 2006; Wada et al. 2005; Frölich et al. 1998), but it was only the STZ-icv rat model which has provided evidence that IRBS forms the core of the sAD pathophysiology by preceding and eventually triggering the sAD-like pathology. This hypothesis seems to gain more and more attention in the sAD research field, in particular now when the "amyloid cascade" hypothesis has been argued and the idea of

To conclude, a growing body of evidence indicates similarities in cognitive deficits, brain neurochemical and structural pathology between the sAD patients and STZ-icv rats which have been proposed as a non-transgenic model of sAD. Considering the recognition of the IRBS condition in the sAD patients postmortem, the development of IRBS following the STZ-icv treatment suggests that the STZ-icv rat model might be a promising experimental tool in both the sAD etiopathogenesis research and novel drug testing. Preliminary data on IRBS being

causally different sAD forms is starting to be considered (Correia et al. 2011).

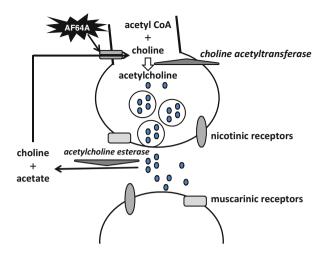
the primary event in this non-transgenic model which could eventually progress and initiate the pathological hallmarks of sAD might provide a new perspective in the research of early diagnosis of this disease in humans. Considering the literature data on environmental and food contaminant exposures to compounds structurally related to STZ, the hypothesis that environmental toxins might play a significant role in the sAD etiopathogenesis should not be neglected (Landrigan et al. 2005; Tong et al. 2009; de la Monte and Tong 2009; de la Monte et al. 2009). In line with that, the STZ-icv rat model provides a wide range of application in the translational neuroscience, and hopefully its further characterization and validation, in particular regarding the long-term follow-up, will bring about the change in elucidating of the sAD enigma.

## 2.3 Streptozotocin-icv + Transgenic Mice

Considering the STZ-icv rat as a model for the sporadic form and the transgenic (Tg) mice overexpressing beta-amyloid protein precursor (APP Tg2576) as a model for the familial (hereditary) form of AD (Hsiao et al. 1996), the question has arisen on how the STZ-icv treatment would affect the Alzheimer-like pathology in the latter model. Generally, in APP Tg2576 mice, a significant increase in soluble A $\beta$ 1-40 and A $\beta$ 1-42/43 peptides is found from 3 to 9-10 months of age, and the senile amyloid plaques are formed at the age of 1 year and older (Hsiao et al. 1996), while the impaired spatial memory is found at the age of 9-10 months and later (Ashe 2001). STZ-icv treatment of 3-month-old APP Tg2576 mice induced increased mortality, and measurements done at the age of 9 months (6 months after the STZ-icv treatment) demonstrated reduced spatial cognition and increased cerebral aggregated A $\beta$  fragments, total tau protein, and congophilic amyloid deposits in comparison to the untreated APP Tg2576 mice (Plaschke et al. 2010b). These changes were associated with decreased GSK- $3\alpha$ /GSK- $3\beta$ ratio (phosphorylated/total) and additionally with a linear negative correlation between Aβ1-42 and cognition and between GSK-3α/GSK-3β ratio and aggregated  $A\beta 1-40 + 1-42$ , indicating that the STZ-icv treatment induced significant aggravation of the Alzheimer-like pathology in this transgenic mice model of familial AD.

### 2.4 Cholinotoxin Model

One of the oldest theories of the AD etiopathogenesis proposed the cholinergic deficits, demonstrated as a dramatic decrease of ChAT in the hippocampus and frontal cortex and the marked reduction in cholinergic neuron counts in the nucleus basalis in AD patient brains post-mortem, to be the core in the AD pathophysiology (Greenwald and Davis 1983). Importantly, ChAT deficit has been shown to correlate with histopathologic changes in the brain and psychologic test scores in AD patients (Greenwald and Davis 1983). In line with that, the putative cholinergic



**Fig. 3** *Mechanism of action of cholinotoxin AF64A in the cholinergic synapse.* Acetylcholine is synthesized in the presynaptic cholinergic neuron from acetyl coenzyme A (acetyl-CoA) and the precursor choline by means of the choline acetyltransferase. Degradation of acetylcholine in the synaptic cleft is mediated by the acetylcholine esterase, generating acetate and choline. The uptake of choline back into the presynaptic nerve ending is mediated by the high-affinity choline membrane transporter, and this process provides the only source of choline necessary for the acetylcholine synthesis. AF64A is an ethylcholine mustard aziridinium ion, a neurotoxic choline analogue, which is taken up by the high-affinity choline transport system into cholinergic neurons (Fisher et al. 1982)

neurotoxin AF64A was injected intracerebroventricularly in order to determine whether it would produce specific damage to the cholinergic cell bodies and thus be useful in developing animal models of AD (Fig. 3). AF64A is an ethylcholine mustard aziridinium ion, a neurotoxic choline analogue, which is taken up by the high-affinity choline transport system into cholinergic neurons (Fisher et al. 1982). Intracerebroventricular administration of 65 nmol of AF64A to mice was not acutely lethal, but produced delayed behavioral effects similar to those found after parenteral administration of AF64A (ataxia, hypokinesia, weight loss) (Fisher et al. 1982). Seven days following the icv treatment, there was a significant decrease in acetylcholine content in the cortex, striatum, and hippocampus, but no change in choline levels. Acetylcholine content was still significantly reduced in the hippocampus at 3 weeks after this treatment (Fisher et al. 1982). The reduction in activity of choline acetyltransferase and high-affinity choline transport paralleled the reduction in acetylcholine measured at 7 days post AF64A treatment, whereas muscarinic receptors in all three brain areas were unchanged (Fisher et al. 1982). Data has indicated that AF64A is a presynaptic chemical neurotoxin, capable of inducing a persistent deficiency in central cholinergic transmission. The work that followed has demonstrated that AF64A-icv injections in addition to causing cholinergic damage in the septum cause also glial lesions in the lateral septal nucleus and in the lateral zones of the hippocampus (Dudas et al. 2004). Small amounts of AF64A (0.01 nmol/ $\mu$ L) injected in each ventricle had little effect on the appearance of the nucleus basalis of Meynert (nbM) or on the ChAT levels in the cortex (Kozlowski and Arbogast 1986). Larger amounts of AF64A (0.02 and 0.05 nmol/µ L and 0.02 nmol/10  $\mu$ L, respectively) produced a loss of diffuse acetylcholinesterase (AChE) staining in the nbM and a loss of large positively staining neurons as well as a significant reduction of ChAT activity in the central portion of the cortex without affecting non-cholinergic neurons (Kozlowski and Arbogast 1986). ChAT activity in the AF64A rat model was preferentially reduced in tissue samples of the dorsal compared to the ventral hippocampus (Ayala-Grosso and Urbina-Paez 1999). At the gene level, AF64A-icv injection (2 nmol) was accompanied by a tenfold increase in the AChE mRNA and a twofold decrease in the butyrylcholinesterase (BChE) and AChE mRNA expression in the hippocampus by day 7 with the latter remaining decreased until day 60 posttreatment (Lev-Lehman et al. 1994). AF64A treatment affects also monoaminergic transmission in the brain, as demonstrated by a dosedependent decrease in noradrenaline level and increased level of its metabolite in the hippocampus 2 days after the treatment (Hörtnagl et al. 1989) and increased serotoninergic neuronal activity and decreased densities of 5-HT uptake sites in different brain regions (Park and Lim 2010).

AF64A-treated rats exhibit learning and memory deficits which have been recorded in different cognitive tests (Nakahara et al. 1988; Lim et al. 2001; Mukhina et al. 2004). In the Morris water maze, these rats showed the delayed latencies to find the platform from the 6th day after the infusion, while in pretrained rats, AF64A treatment caused the significant delay of latency at the 7th day, but not the 8th day (Lim et al. 2001). In the passive avoidance test, AF64A-treated rats shortened the latency 1.5 h but not 24 h after the electroshock, suggesting that AF64A impairs the learning and memory but also that the disturbed memory might rapidly recover after the first retrain (Lim et al. 2001). No literature data are available regarding the amyloid and tau protein pathologies as well as the IRBS in the cholinotoxin AF64A animal model of AD which has not been frequently used in the contemporary AD research.

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# Fusion Models and "Fusioning" in Parkinsonism: Protection and Restoration by Exercise

Trevor Archer and Anders Fredriksson

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#### Abstract

Fusion models, or the "fusioning" of models, of Parkinson's disease (PD) to attain the sufficiency of disorder etiopathogenesis have spurred the development of laboratory models incorporating neurotoxin treatments in combination with genetic manipulations of animals studied. The present review describes fusion from two directions: (i) through the fusioning of neurotoxin and genetic models, and, It was observed that wheel-running exercise (four 30-min sessions/week) attenuated the motor deficits, both assessed as distance run in the running wheels and as locomotor, rearing and total activity counts in the motor activity test chambers, and the dopamine (DA) loss induced by the DA neurotoxin, MPTP compared with the no exercise MPTP group (one 30-min session/week).

### Keywords

DA • Exercise • Mice • Motor activity • MPTP • PD

T. Archer (🖂)

A. Fredriksson

Department of Neuroscience, Psychiatry, Ulleråker, University Hospital, Uppsala, Sweden

Department of Psychology, University of Gothenburg, Gothenburg, Sweden e-mail: trevor.archer@psy.gu.se

## 1 Introduction

Parkinson's disease (PD) is second only to Alzheimer's disease as a neurodegenerative disorder afflicting humankind with four to five million individuals, diagnosed or undiagnosed in the USA alone (Korell and Tanner 2005). PD most commonly affects individuals in their sixth decade or later and involves several risk factors including pesticides, head trauma, and aging (Allam et al. 2005). Pathways mediated through mitochondrial actions are implicated heavily in the neurodegenerative cascades that result in dopaminergic cell death (Jordan et al. 2003, 2011), with mitochondrial outer membrane permeabilization intimately associated with the apoptotic process (Galluzzi et al. 2009; Gomez-Lazaro et al. 2008). PD is a relatively common, idiopathic neurodegenerative movement disorder with the cardinal features of impaired motor function, resting tremor, rigidity, akinesia/bradykinesia, and postural instability (Gaggelli et al. 2006; Jankovic 2008; Lees et al. 2009). It is a progressive neurodegenerative disorder with late onset and, compared with familial forms, is associated most often with advanced age (>55 years of age). PD symptoms are manifested following 50–75 % loss of striatal dopamine (DA, Riederer and Wuketich 1976; Scherman et al. 1989) and 30-60 % destruction of nigrostriatal DA neurons (Cheng et al. 2010; Fearnley and Lees 1991). Axonal pathology in PD is supported by various sources, not least genetic mutational models (Li et al. 2009; Mortibovs et al. 2008; Saha et al. 2004). For example, Mortiboys et al. (2010) obtained impaired mitochondrial function and morphology in LRRK2 (G2019S)-mutant PD patient tissue, i.e., shared features in early-onset and late-onset PD. The consensus of these and other studies indicating alterations of axonal transport (Meissner et al. 2011; Weihofen et al. 2009) reinforces the notion that neurons in the substantia nigra degenerate through "retrogradely perishing" axonopathy whereby deterioration originates in the distal axon and proceeds to the cell body temporally (Antenor-Dorsey and O'Malley 2012). Converging pathogenic mechanisms imply a restricted number of mechanisms that include oxidative stress, mitochondrial defects and dysfunctionality, proteolytic stress, and neuroinflammation (Braak et al. 2006).

The pathophysiology of PD involves dopaminergic neuron death and accumulation of Lewy bodies associated with mutations in  $\alpha$ -synuclein, a 14-kDa protein predominantly expressed in the brain and CNS (Rasia et al. 2005). PD patients show decreased levels of presynaptic DA neuron terminal markers in the basal ganglia (Felicio et al. 2009), consistent with loss of dopaminergic terminals due to degeneration of neuronal cell bodies in the substantia nigra pars compacta (Hattori et al. 2006). They also exhibit decreased levels of DA transporters (DATs) and vesicular monoamine transporter type 2 (VMAT2) as well as reduced activity of dopa decarboxylase, assessed by striatal conversion of L-Dopa to DA, according to PET and SPECT analyses. Wu et al. (2012) using MRI showed that the substantia nigra pars compacta expressed a decreased connectivity with several regions, including the striatum, globus pallidus, subthalamic nucleus, thalamus, supplementary motor area, dorsolateral prefrontal cortex, insula, default mode network, temporal lobe, cerebellum, and pons in patients compared to controls. They found that L-Dopa administration partially normalized the pattern of connectivity to a similarity such as that expressed by the healthy volunteers involving causal connectivity of basal ganglia networks from the substantia nigra pars compacta. Postsynaptic  $D_2$  DA receptors ( $D_2R_s$ ) are either unaffected or increased in the striatum of untreated PD patients (Antonini et al. 1994). Oxidative injury appears to be one effect of  $\alpha$ -synuclein ( $\alpha$ -Syn) aggregates and could ultimately produce neuronal cell death.  $\alpha$ -Syn, a 140 residue, intrinsically disordered protein is localized in presynaptic terminals of DA neurons (Yang et al. 2010). Autonomic nervous system involvement occurs at early stages in both PD and incidental Lewy body disease and affects the sympathetic, parasympathetic, and enteric nervous systems. It has been proposed that  $\alpha$ -Syn pathology in PD has a distal to proximal progression along autonomic pathways. According to Braakian notions, the enteric nervous systems is affected before the dorsal motor nucleus of the vagus, and distal axons of cardiac sympathetic nerves degenerate before there is loss of paravertebral sympathetic ganglion neurons. Cersosimo and Benarroch (2012a) have shown that consistent with neuropathological findings, some autonomic manifestations such as constipation or impaired cardiac uptake of norepinephrine precursors occur at early stages of the disease even before the onset of motor symptoms (cf. Braak et al. 2006; Cersosimo and Benarroch 2012b; Hawkes et al. 2007).

## 2 Fusion in PD

The notion concerning the "fusion of models" of PD, the current and latest phase of Parkinsonian model development, has provided a major basis for studying the etiopathogenesis of the disorder through the presentation of a whole new set of animal models and clinical scenarios whereby gene x environment interactions may be analyzed (Manning-Bog and Langston 2007). Several models of PD induced by treatment with neurotoxins included 1-methyl-4-phenyl-1,2,3,6systemic tetrahydropyridine (MPTP), the pesticide rotenone, the herbicide paraquat, and liposaccharide (LPS) occurring in outer membrane of gram-negative bacteria, which combine to provide dopaminergic vulnerability for PD that endows the "multiple-hit" aspect to establish PD symptom profiles, if not a complete pathophysiology. Other models of PD are based upon some form of genetic mutation of  $\alpha$ -synuclein or Parkin deficiency, DJ-1 deficiency, and *PINK-1* and *LRRK2* genes that are employed in transgenic models (Manning-Bog and Langston 2007). Remarkably, Westerlund et al. (2010) advocated genetic factors as the major causative factor in PD due to the multiplicity of genes now implicated in the etiopathogenesis that involve several molecular pathways and downstream effectors affecting the trophic support and/or the survival of DA neurons. Nevertheless, much evidence suggests that multiple factors combining environmental agents and genetic predispositions orchestrate the fate of DA neurons (de Lau and Breteler 2006; Gupta et al. 2008; Schapiro et al. 2009). Manning-Bog and Langston (2007), as well as others (Nieto et al. 2006; Song et al. 2004), have presented the notion of "fusion" between transgenic mouse lines, which indicate subtle alterations with risk factor relevance but generally fail to show overt damage to the nigrostriatal DA system, and laboratory models of neurotoxin-induced DA depletion and altered neurotransmission. Several instances of genetic and neurotoxin model have been reported: Liu et al. (2012) have found that SIRT2 (a sirtuin) deacetylates Foxo3a, increases RNA and protein levels of Bim, and, as a result, enhances apoptosis following MPTP treatment and that chronic MPTP- induced neurodegeneration was prevented by the genetic deletion of SIRT2 in mice. The deletion of SIRT2 led to the reduction of apoptosis due to an increase in acetvlation of Foxo3a and a decrease in Bim levels. Peroxisome proliferator-activated receptor-gamma coactivator- $1\alpha$  (PGC- $1\alpha$ ), a transcriptional coactivator regulating cell metabolism, mitochondrial biogenesis, oxidative stress, and gene expression, modulates the sensitivity of neurons in the brain to insults; PGC-1a knockout mice were shown to be supersensitive to MPTP (St-Pierre et al. 2006). Several recent studies have shown that PGC-1 $\alpha$  is necessary for the integrity of DA neurons through its actions against oxidative stress and enhancing neuronal viability (Breidert et al. 2002; Zheng et al. 2010). Mudo et al. (2012) applied resveratrol to modulate/activate PGC-1a in dopaminergic SN4741 cells in vitro via the deacetylase SIRT1, thereby enhancing PGC-1 $\alpha$  gene transcription with elevated SOD2 and Trx2.

The necessity for a model fusion approach has been spurred by several avenues of discontent stemming from what has been described as a "myopic" predilection for neurotoxin-based assaults of the dopaminergic nigrostriatal system (Langston 2006); all viable notions pertaining to the etiopathogenesis of the disorder must accommodate the circumstance that PD presents not a single entity that is the cumulative effect of a dopaminergic structural and functional defect accompanied by several related deficits but expresses rather a concatenation of destructive regional, cellular, and molecular events brought together by a combination of genetic and environmental factors (Litvan et al. 2007a, b). The existence of familial (i.e., genetic) forms of the disorder that appears to account for about 10–18 % of PD diagnoses has been considered (Wirdefeldt et al. 2011). Gao et al. (2012) showed that a family history of Parkinsonism was associated with PD risk (odds ratio = 2.71, 95 % confidence interval, 1.97-3.74), but at the same time was affected marginally, or not at all, by further adjustment for these SNPs and environmental exposures; they obtained a significant interaction with rs2896905 at SLC2A13, near LRRK2, with 32-35 % higher risk levels. Blandini and Armentero (2012) have classified animal models of PD on the basis of systemic/ intracerebral administration of neurotoxins that replicated pathological and phenotypic characteristics of the disorder yet describe the proliferation of mammalian and nonmammalian transgenic models presenting disease-inducing mutations identified for monogenic forms of familial PD. Nevertheless, the current trend towards "fusioning" of animal models in this respect holds both specific characteristics that may not be commensurate with essential aspects of the disorders, such as "staging" (Archer et al. 2011c), or limitations requiring much careful consideration prior to selection and application. Soreq et al. (2012) carried out a meta-analysis on 131 brain region transcriptomes from mice overexpressing native or mutated  $\alpha$ -synuclein (SNCA) with or without the protective HSP70 chaperone or exposed MPTP, with or without the protective acetylcholinesterase variant, AChE-R, all of which presented risk-inducible and protection-suppressible transcript modifications. Their conclusions imply that metal ions mediated the "cross talk" between nuclear and mitochondrial pathways by both environmental and genetic risk and those protective factors which eventually culminate in neuroinflammation in the PD cascade. In nonhuman primate studies of PD, both the motor and cognitive capacities of nonhuman primates (NHPs) provide levels of structural and functional complexity that permits laboratory observations to attain insights for applications in the clinical realities (Capitanio and Emborg 2007).

Applying a different point of departure to the investigation of Parkinsonism in the laboratory, fusion dynamics (mitochondrial biogenesis, bioenergetics, dynamics, transport, quality control, and the genes involved) are critical for the life cycle and maintenance of dopaminergic neuron mitochondria. Mitochondrial integrity, dependent upon specific morphological, biochemical, and physiological features, is of critical importance for the maintenance and survival of neurons (Exner et al. 2012) with dysfunctions linked to severe cellular consequences that result in neurodegenerative diseases and aging. Thus, Arduino et al. (2011) have proposed that the cellular demise and neurodegenerative propensity in PD pathophysiology may be modulated by an interaction between mitochondrial dysfunction, mitochondrial trafficking disruption, and impaired autophagic clearance. Sporadic PD case studies, neurotoxin-mediated animal models, and genetically based (familial PD) PD etiology imply that mitochondrial dysfunction/ dysregulation constitutes an earl feature of the disorder (Burman et al. 2012; Cooper et al. 2012; Moran et al. 2012). Increased mitochondrial damage due to complex I inhibition, derangement of Ca<sup>++</sup>, oxidant stress, or impaired clearance of mitochondrial debris leads to the accumulation of nonfunctional organelles contributing to neuronal disrepair (McCoy and Cookson 2012). Several gene mutations regulate the responses to cellular stress with enhanced sensitivity to mitochondrial neurotoxins and oxidative stress (Alberio et al. 2012; Heo et al. 2012; Rochet et al. 2012). Macroautophagy is associated with prevention and limitation of mitochondrial damage and debris through degradation of superfluous or severely damaged mitochondria (Lynch-Day et al. 2012; Wang and Klionsky 2011). Mitochondrial fusion processes, distinguished from membrane fusion mechanisms in secretory pathway or membrane-bound organelles, are divided into three events: "docking," outer membrane fusion, and inner membrane fusion. Injuries to mitochondria impair fusion activate fission-dependent fragmentation and trigger mitophagy (Kim et al. 2007). For instance, mitochondria-derived reactive oxygen species (ROS) at low concentrations may act as signaling molecules and induce mitophagy (Scherz-Shouval et al. 2007). Mitochondrial quantity and quality are regulated to prevent excess energy and harmful ROS. Mitophagy limits the mitochondria to a basal level to fulfill cellular energy requirements and preventing excess ROS production (Kurihara et al. 2012). Arduino et al. (2011) have proposed that cellular demise and neurodegeneration in PD stem from the interactions between mitochondrial dysfunction and trafficking disruption and autophagic clearance. In this context, fusion-fission models of the disorder pertain to conditions wherein mitochondrial structural-functional integrity through the effective working of cellular autophagy and mitophagy has been compromised (Santos and Cardoso 2012; Youle and van der Bliek 2012).

From a PD perspective, axonal integrity and function are disrupted also by a variety of environmental and laboratory neurotoxins, including 6-hydroxydopamine (Segura Aguilar and Kostrzewa 2004; Kostrzewa et al. 2006), MPTP (see below, Annese et al. 2012; Villalba and Smith 2011), rotenone (Norazit et al. 2010; Press and Milbrandt 2008), and paraquat (Brooks et al. 1999; Peng et al. 2004). Both MPP<sup>+</sup> and rotenone disrupt axonal integrity and function by inhibiting mitochondrial complex I activity (Sterky et al. 2012) and depolymerizing microtubules leading to axon fragmentation and decreased synaptic function (Cabeza-Arvelaiz and Schiestl 2012; Cartelli et al. 2010; Hongo et al. 2012; Qureshi and Paudel 2011); the neurotoxic insults of these agents linked to inhibition of axon transport directly (Behrouz et al. 2007; Kim-Han et al. 2011; Morfini et al. 2007). Axon degeneration, characteristic to neurodegeneration, shares many morphological features with events expressed in acute injuries; the expression of the Wallerian degeneration slow  $(Wld^{S})$  transgene delays nerve degeneration in both types of insult (Wang et al. 2012). Wallerian degeneration occurs through crushing/cutting nerve fibers whereby the section of the axon separated from the cell body degenerates distal to the injury; Wallerian-type degeneration occurs in PD through compromise of axonal transport. The Wallerian degeneration slow fusion protein,  $Wld^{S}$ , delays axonal degeneration about 10-fold (Avery et al. 2012; Chitnis et al. 2007; Coleman and Freeman 2010; Samsam et al. 2003; Takada et al. 2011) and blocks axon degeneration in several models, such as animal models of PD (Mi et al. 2005; Sajadi et al. 2004) that include the MPTP model (Antenor-Dorsey and O'Malley 2012; Hasbani and O'Malley 2006). Using mutant mice and lentiviral transduction of DA neurons, Antenor-Dorsey and O'Malley (2012) showed that *Wld<sup>S</sup>*, but not Nmnat1, Nmnat3, or cytoplasmically targeted Nmnat1, protected DA axons from MPTP treatment that resulted in MPP<sup>+</sup>. Lentiviral transduction refers to a form of "gene therapy" used to designate a process through which viruses are exploited to carry specific genetic information into a cell, thereby transducing the cells' function (cf. Tolu et al. 2010). NAD<sup>+</sup> synthesis was not required but was, by itself, sufficiently axonally protective and was additive in effect together with  $Wld^{S}$ . It appears that  $Wld^{S}$  and NAD<sup>+</sup> act through separate pathways to protect DA neurons through preserving mitochondrial health and/or maintaining cellular metabolism.

The regulation of processes incorporating mitochondrial dynamics encompasses essential mechanisms that control the eventual fate of cells, such as dopaminergic neurons (Liesa et al. 2009); these processes include fission, fusion, and mitophagy, with fusion and fission acting antagonistically, and imbalances in mitochondrial dynamics contributing to PD (Cho et al. 2010; Lees et al. 2009; Su et al. 2010). Mitophagy eliminates damaged or aged mitochondria, thereby maintaining mitochondrial quality, an autophagy-dependent degradation of mitochondria that contributes to the removal of dysfunctional, aged, or excess mitochondria (Hirota et al. 2012). Galindo et al. (2012) have shown that the regulation of dynamic mitochondrial processes such as fusion, fission, and mitophagy has been unfolded as cascades contributing to important mechanism controlling neuronal fate.

### 3 MPTP Model of PD

Repeated administration of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) to C57/BL6 mice produces selective lesions of dopaminergic neurons in substantia nigra pars compacta and long-lasting depletion of dopamine (DA) in basal ganglia (Jackson-Lewis et al. 1995; Jones-Humble et al. 1994; Langston 1985; Langston et al. 1984). The susceptibility of mice to the neurotoxic actions of MPTP can be quite variable, depending on gender and strain differences, expressed in functional, neurochemical, and histochemical analyses (Schwarting et al. 1999; Sedelis et al. 2000a, b, 2001, 2003). C57/BL6 and Swiss Webster strains were shown to differ in c-Jun N-terminal kinases (JNKs) and c-JUN activation in response to MPTP. JNKs, of the mitogen-activated protein kinase family, are responsive to stress stimuli, such as cytokines, ultraviolet irradiation, heat shock, and osmotic shock; c-Jun is the name of a gene and protein that, in combination with c-Fos, forms the AP-1 earlyresponse transcription factor. MPTP induced COX-2, an enzyme responsible for inflammation and pain, responding exclusively in C57/BL6 mice (Boyd et al. 2007). MPTP, administered systemically, induces Parkinsonism in human and nonhuman primates (Langston 1985) that results in the loss of substantia nigra cells in the pars compacta of adult animals (Chiueh et al. 1985). MPTP selectively destroys nigrostriatal neurons, thereby inducing acute, subacute, long-lasting, and even permanent effects that resemble certain features of PD, particularly the hypokinesia effects (Schultz et al. 1989). Systemic administration of MPTP (2  $\times$ 40 mg/kg, s.c.) caused L-Dopa reversible hypoactivity (Fredriksson et al. 1990; Sundström et al. 1990). A less rigorous dose regime, e.g.,  $2 \times 20$  or 25 or 30 mg/kg, of MPTP has been found not to reduce motility in the C57 black mice, although DA brain concentrations may indicate up to 50–80 % reductions (Heikkila et al. 1989; Sonsalla and Heikkila 1986), unless given much more repeatedly (cf. Kurz et al. 2007). The parameters of MPTP treatment neurotoxicity in mice are long-lasting (up to and beyond 52 weeks after treatment), with strong correlations between the functional deficits, particularly hypokinesia, the main biomarker, severe DA depletions, and a dose- and time-dependent recovery of several parameters of motor behavior following treatment with the DA precursor, L-Dopa (Archer and Fredriksson 2003; Fredriksson and Archer 1994; Fredriksson et al. 1999).

## 4 Exercise Intervention in PD

In the unilateral 6-hydroxydopamine rat model of PD, Tillerson et al. (2001) abolished the lesion-induced motor asymmetry by forcing rats to use the affected (contralateral)

limb, whereas forced nonuse exacerbated the injury (Tillerson et al. 2002). Both DA and 3,4-dihydroxyphenylacetic acid (Dopac) were elevated markedly in "casted" 6-OHDA-treated rats (forced to use the contralateral limb) compared with "non-casted" rats (Cohen et al. 2003). More recently, Choe et al. (2012) showed that exercise recovers PD-induced dopaminergic neuron loss and contralateral soleus muscle atrophy. Using the DA neurotoxin, MPTP as their model of PD, Sung et al. (2012) observed that treadmill exercise prevented dopaminergic neuronal loss by inhibiting brain inflammation through suppression of microglial activation in the PD mice. Archer and Fredriksson (2010) found that daily running-wheel activity attenuated the hypokinesia effects of MPTP in both a concentrated ( $2 \times 40$  mg/kg, 24-h interval) and progressive  $(1 \times 40 \text{ mg/kg}, \text{weekly doses over 4 weeks})$  schedule with regard to spontaneous motor behavior and activity following a subthreshold dose of L-Dopa. The loss of DA in each case was attenuated by exercise also (experiment I, 61 % of control vs. 17 %; experiment II, 24 % vs. 11 %). Using the progressive schedule of MPTP treatment and extending the exercise intervention from 7 to 14 weeks, it was shown that spontaneous motor activity after MPTP was close to restoration, whereas activity after subthreshold L-Dopa was completely recovered (Fredriksson et al. 2011); DA levels were restored from 17 % (non-exercised) to 64 % in the 14-week exercise intervention, and levels of brain-derived neurotrophic factor were increased significantly. Also, both the functional and DA deficits by MPTP were attenuated even by delayed introduction of exercise (Archer and Fredriksson 2012).

Zigmond et al. (e.g., Cohen et al. 2003; Tillerson et al. 2001, 2002) have utilized the unilateral 6-hydroxydopamine (6-OHDA) rat model of PD whereby they increased the use of the normally denervated contralateral by constraining the other forelimb with a cast, thereby reducing the function and neurochemical effects of the DA neurotoxin evidenced by marked elevations of DA and metabolites. Using MPTP to deplete DA, Klaissle et al. (2012) analyzed newborn cells in the substantia nigra at different stages of maturation and time points as a function of voluntary physical activity, environmental enrichment, and L-Dopa treatment. They observed a motor activity-induced, running-wheel increase in neuroglial antigen-2-positive cells and mature oligodendrocytes in the substantia nigra of healthy mice. In MPTP-treated mice, the stimulation of neuroglial antigen-2-positive cell generation and oligodendrogenesis was dependent upon L-Dopa administration. The authors have suggested that DA is the key regulator for generation of neuroglial antigen-2-positive cells and oligodendrocytes in the substantia nigra that ought to be taken into account as an important contributor to endogenous nigral cellular plasticity. The integrity of the nigrostriatal system under MPTP insult is linked further with dynamics of glial cell activation: CD11b expression, a surface marker on the plasma membrane of microglia and the  $\alpha$ -component of several integrins, is increased in the striatum and substantia nigra following MPTP (Kang et al. 2008; Singh et al. 2011). Integrins, transmembrane receptors that mediate the attachment between neurons and surrounding tissues, pass information about the chemical composition and mechanical status of the extracellular matrix. Sung et al. (2012) have shown that treadmill running exercise prevented the loss of dopaminergic neurons and ameliorated coordination dysfunction in MPTP mice by inhibiting brain inflammation through microglial activation. The effects of physical exercise upon functional and biomaker integrity have been amply shown (Archer 2011, 2012; Archer et al. 2011a, b, 2012; Archer and Kostrzewa 2012; Archer and Fredriksson 2013).

## 5 Conclusion

The fusion modeling of PD, whether viewed as neurotoxin-genetic model fusions or fusioning within mitochondria, offers an abundance of structural, epigenetic, bio-, and even neurophysiologic markers (Baziyan 2012) as evidence for essential neurodegenerative processes involved in the disorder, if not a complete etiopathogenesis. There is an emerging consensus that genetic- and toxin-induced animal together with cellular models and postmortem human brain tissue provide evidence that mitochondrial dysfunction/dysregulation forms the basis disorder pathophysiology (Chaturvedi and Beal 2012; Van Laar and Berman 2012). From the distinguishing perspective of neuroprotection and/or neurorestoration, Ohshima-Hosoyama et al. (2012) have combined the human insulin receptor (HIR) with glial-derived neurotrophic factor (GDNF) in the fusion protein, HIRMAb-GDNF, in order to induce neuroprotection in Parkinsonian monkeys. However, their results demonstrated that HIRmAb-GDNF dosing in a monkey model of PD was not an effective neuroprotective strategy. Physical exercise has proven effective even using other laboratory models of PD: using the chronic haloperidol administration model of PD, Baptista et al. (2013) showed that exercise alleviated haloperidol-induced akinesia (see also Abrantes et al. 2012). Finally, taking into account noninvasive, non-pharmacological strategies to prevent/restore against brain degeneration in PD, physical exercise has been shown to offer a remarkably effective strategy, alleviating and even restituting brain tissue and mitochondrial dysfunction. Marquis-Aleixo et al. (2012) have described the role of physical exercise in the modulation of the mechanisms involved in neuroprotection including the activation of signaling pathways underlying brain protection.

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# Iron-Induced Dopaminergic Cell Death In Vivo as a Model of Parkinson's Disease

Manfred Gerlach, Kay L. Double, and Peter Riederer

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#### Abstract

Histopathological, biochemical, and in vivo brain imaging techniques have revealed a consistent increase of total iron in the substantia nigra pars compacta (SNc) of Parkinson's disease (PD) patients. Interestingly, the increased iron was shown to be restricted to the SNc and occurs in the ferric ( $Fe^{3+}$ ), rather than ferrous ( $Fe^{2+}$ ), form of the metal.

M. Gerlach (🖂)

K.L. Double

#### P. Riederer

University of Würzburg, Department of Child and Adolescent Psychiatry, Psychosomatics and Psychotherapy, Laboratory for Clinical Neurobiology, Würzburg, Germany e-mail: manfred.gerlach@uni-wuerzburg.de

The University of Sydney, Discipline of Biomedical Science, School of Medical Science, Sydney Medical School, Sydney, Australia e-mail: kay.double@sydney.edu.au

University of Würzburg, Department of Psychiatry, Psychosomatics and Psychotherapy, and National Parkinson Foundation Centers of Excellence for Neurodegenerative Diseases Research, Laboratory for Clinical Neurochemistry, Würzburg, Germany e-mail: peter.riederer@mail.uni-wuerzburg.de

The objective of this chapter is to discuss the evidence for dopaminergic cell death in the rat following unilateral ferric iron injections into the SN. In addition, this chapter will briefly review findings showing that NM-bound ferric iron represents a pool of iron that, under certain circumstances, can be released to interact with free radical-producing pathways and the ubiquitin proteasome system, ultimately leading to dopaminergic nerve cell death in the rat. Special attention will be paid to summarize evidence that iron-induced dopaminergic cell death reproduces key features of PD and mimics molecular mechanisms underlying dopaminergic cell death in PD.

#### **Keywords**

Animal models • Iron • Neuromelanin • Oxidative stress • Parkinson's disease • Progression

#### 1 Introduction

Parkinson's disease (PD) is the most common neurodegenerative movement disorder. The most striking neuropathological characteristic of PD is the relatively specific loss of the pigmented dopaminergic neurons of the substantia nigra (SN) pars compacta (SNc) and the resulting pallor of the midbrain, as well as the development of abnormal  $\alpha$ -synuclein-immunopositive inclusion bodies within the boundaries of the pigment (Jellinger 2012). While the full clinical syndrome of PD is a complex multi-transmitter disorder, the characteristic motor symptoms associated with the sporadic, and all genetic, forms of this disorder result from the degeneration of the dopamine-producing neurons of the SNc. While the trigger for this relatively selective neuronal vulnerability remains unknown, the cascade of degenerative events leading to cell death is beginning to be understood. The major hypotheses believed to contribute to the eventual demise of nigral dopamineproducing cells include altered protein handling, disturbed iron homeostasis, oxidative stress, mitochondrial dysfunction, and neuroinflammation (Gerlach et al. 1994, 2006a; Sian-Hulsmann et al. 2011; Schapira and Jenner 2011). Age is the single most important risk factor for PD, and the biochemical changes that are consequences of aging amplify these abnormalities in the PD brain (Schapira and Jenner 2011).

Histopathological, biochemical, and in vivo brain imaging techniques have revealed a consistent increase in total SNc iron in PD (Gerlach et al. 1994, 2006a; Sian-Hulsmann et al. 2011). Interestingly, the increased iron was shown to be restricted to the SNc and occurs in the ferric ( $Fe^{3+}$ ), rather than ferrous ( $Fe^{2+}$ ), form of the metal (Sofic et al. 1988).

Iron increases have also been observed in the brain of nonhuman primates and rodents after exposure to dopaminergic neurotoxins such as 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP; Mochizuki et al. 1994), 6-hydroxydopamine (6-OHDA; Oestreicher et al. 1994), methamphetamine (Melega et al. 2007), and lipopolysaccharide (Zhang et al. 2012). For a detailed treatment of MPTP,

6-OHDA, and methamphetamine, the interested reader is referred to corresponding chapters in this handbook.

Increased iron deposits in the SNc in PD may result from several sources, including a damaged or "leaky" blood-brain barrier in the SN and disturbances of iron uptake, storage, and transport as neurodegeneration progresses (Gerlach et al. 2006b; Sian-Hulsmann et al. 2011). Major iron stores are ferritin and hemosiderin in glial cells, and neuromelanin (NM) in neurons. Age- and disease-dependent overload of iron storage molecules may result in iron release upon reduction. Consequently, the low molecular weight chelatable iron complexes may trigger redox reactions (e.g., oxidative stress), leading to damage of biomolecules, ultimately leading to dopaminergic nerve cell death. This damage includes a defect in complex I in mitochondria, DNA fragmentation, lipid peroxidation of cell membranes, protein misfolding. and disturbance of calcium homeostasis. Additionally, upon neurodegeneration, there is strong microglial activation that can be another source of high iron concentrations in the brain. Although the current evidence suggests that increased brain iron may be a secondary result of neuronal degeneration and reactive gliosis in several neurological disorders affecting the basal ganglia including PD, progressive supranuclear palsy, and multisystem atrophy, the question of whether iron-associated degenerative pathways are a significant factor driving progressive neuronal death in these disorders is yet to be definitively answered.

The objective of this chapter is to discuss the evidence for dopaminergic cell death in the rat following unilateral ferric iron injections into the SN. In addition, this chapter will briefly review findings showing that NM-bound ferric iron represents a pool of iron that, under certain circumstances, can be released to interact with free radical-producing pathways and the ubiquitin proteasome system, ultimately leading to dopaminergic nerve cell death in the rat. Special attention will be paid to summarize evidence that iron-induced dopaminergic cell death reproduces key features of PD and mimics molecular mechanisms underlying dopaminergic cell death in PD.

# 2 Effects of Unilateral Intranigral Injections of Ferric Iron

The unilateral injection of 50  $\mu$ g of ferric iron into the right SN of the rat produced a strongly altered motor behavior in the animals, which even 3 weeks after application is expressed by a reduction in spontaneous locomotor activity in a strange environment, by a lower frequency of getting up onto the hind-legs, by a transient appearance of "freezing" phenomena and by spontaneous ipsilateral rotations that are accentuated by amphetamine (Ben-Shachar et al. 1991). This altered motor behavior was accompanied by an average 95 % reduction of the dopamine concentration in the ipsilateral striatum, as well as smaller reductions in the dopamine metabolites 3,4-dihydroxyphenylacetic acid (DOPAC) and homovanillic acid (HVA) (85 % or 45 % respectively) measured *post mortem*. Similar changes were also observed at lower iron concentrations (70–350 ng; Sengstock et al. 1992, 1993). In contrast, other neurochemical striatal markers, such as noradrenaline, serotonin, and 5-hydroxyindoleacetic acid, were unchanged (Ben-Shachar et al. 1991; Arendash et al. 1993).

Histopathological investigations predominantly showed damage of the SNc (Sengstock et al. 1992; Arendash et al. 1993). As early as 24 h after intranigral application of 70 ng of ferric iron, iron-stained astrocytes and microglia were demonstrated. Immediately adjacent to these iron-stained cells, there was a severe loss of neurons and a reactive gliosis, still demonstrable after at least 6 months, as well as a decreased number of neurons stained immunohistochemically of tyrosine hydroxylase (TH).

This damage to the dopaminergic system produced by intranigral ferric iron injection was also demonstrated in vivo by voltammetric procedures measuring dopamine release (Junxia et al. 2003) and extracellular DOPAC concentrations (Wesemann et al. 1993, 1994). In the striatum of rats that have been lesioned with 50-µg ferric iron, the DOPAC signal was diminished. It is interesting to note the time course of this signal: During the 1st week following the lesion, there has been a 79 % reduction on the side of the injection compared to the opposite side, while 3–6 weeks later, the signals were diminished by 86 % or 97 % (Wesemann et al. 1994). This was the first time that a toxic insult has been shown to result in a chronic and progressive course in an animal model of PD. This chronic and progressive course provoked by the nigral iron injection was confirmed by other parameters (Sengstock et al. 1994): First, there was a similarly progressive course on dopamine and HVA changes in the striatum, determined at *post mortem*; second, a progressive atrophy of the SN has been found; and third, there was progressive alteration in apomorphine-elicited rotational behavior.

Interestingly, the unilateral injection of low amounts of ferric iron  $(1.5 \ \mu g)$  into the left ventral striatum of the rat induced also a long-lasting disturbance of the dopaminergic function (Kolasiewicz et al. 1995). Ten weeks following injection, there was a decrease of the spontaneous night locomotor activity; 14 weeks post-injection, there was a decrease of spontaneous locomotor activity during the adaptation phase in a new environment and an induction of contralateral rotation behavior following subcutaneous administration of 2 mg/kg apomorphine.

The mechanism of the neurotoxic effect of injected ferric iron has not been extensively investigated and is only partly understood. It is assumed that ferric iron is taken up into the neurons and glia by transferrin-receptors (Arendash et al. 1993). Indeed, it was demonstrated in SH-SY5Y neuroblastoma cells that exposure to ferric iron  $(1-80 \ \mu\text{M})$  leads to a concentration-dependent accumulation of intracellular ferric iron (Nunez et al. 2004). Ferric iron is probably reduced to ferrous iron in the cytosol of these cells. Ferrous iron showed a cytotoxic action in primary cultures of neurons derived from the mesencephalon of rat embryos (Michel et al. 1992). In a reaction similar to the Fenton reaction, excess ferrous iron could catalyze the formation of hydroxyl free radicals from hydrogen peroxide which is an endogenous product of enzymatic dopamine metabolism. This proposed mechanism for the neurotoxic effect was indirectly confirmed by the demonstration of increased amounts of thiobarbituric acid-reactive compounds (a measure of lipid peroxidation) in the SN of ferric iron-infused rats (Arendash et al. 1993;

Wesemann et al. 1993; Sengstock et al. 1997) and the inhibitory effect of desferrioxamine, an iron chelator (Wesemann et al. 1993; Lan and Jiang, 1997). In addition, cell culture studies have demonstrated that exposure of ferric iron (40–150  $\mu$ M) leads to oxidative stress, assessed by reduction of reduced glutathione (GSH) and total glutathione (GSH and GSSG) concentrations (Nunez et al. 2004) as well as activation of redox-sensitive signals (Salvador and Oteiza 2011), and apoptotic cell death (Salvador and Oteiza 2011).

It has also been shown that ferrous iron led to the formation of large perinuclear inclusion bodies in SH-SY5Y cells that overexpress  $\alpha$ -synuclein (Hasegawa et al. 2004). However, the presence of Lewy body-like nigral inclusions has not been established in animals injected with ferric iron into the SN. This is probably the result of the acute administration of iron. It has been observed that inclusions form only in MPTP-treated mice when the neurotoxin is administered continuously at low doses (Fornai et al. 2005).

Extracellular iron has also been shown to increase activated microglia, the principal immune cells in the CNS, and subsequent neurodegeneration in animal models of PD (Shoham and Youdim 2000). Interestingly, an increased SN iron content and increased amounts of activated microglia are associated with SN hyperechogenicity in postmortem brains, whereas increased SN neuromelanin content is related to reduced SN echogenicity (Berg et al. 2010).

# 3 Effects of Unilateral Injections of Human Neuromelanin-Bound Ferric Iron into the Substantia Nigra

NM is considered as an endogenous iron-binding molecule in pigmented neurons as ferritin is primarily located in glia, rather than neurons, and is believed to play a physiological role in intraneuronal iron homeostasis (Double et al. 2000; Gerlach et al. 2006a). Isolated human NM consists of 2.8 % iron as estimated by Mössbauer spectroscopy (Gerlach et al. 1995), while the concentration of ferric iron in the SN has been determined using electron paramagnetic resonance at 11.3-µg iron/mg isolated NM (Shima et al. 1997). Our Mössbauer spectroscopy data demonstrated also that the iron bound to NM is purely ferric iron that is bound to ferritin-like oxyhydroxide clusters (Gerlach et al. 1995). In PD, where nigral iron levels are increased, saturation of high affinity iron-binding sites on NM may overwhelm the protective capacity of this molecule, leading instead to an increase in redox-active iron, and subsequent cellular damage (Gerlach et al. 2008). Support for this theory comes from changes in NM in the PD brain where significantly less iron is bound to NM than that seen in the normal brain (Lopiano et al. 1999). This suggests that changes in iron-binding to NM may result in increased levels of intraneuronal free iron and subsequent cell damage.

Indeed, it has been demonstrated in adult male Sprague-Dawley rats that injection of 1  $\mu$ l of a NM/ferric iron (2.5 nmol or 0.139  $\mu$ g iron) suspension into the ventrolateral region of the left SNc leads to a 50 % reduction of the dopamine neuron number (counted as TH-immunopositive neurons) 8 weeks after injection

compared to that in vehicle-injected animals (Double et al. 2003). The absence of motoric disturbances or a striatal dopamine deficit in these animals suggests a subclinical dopaminergic lesion. This is in contrast to many of the commonly used animal models of PD including the unilateral injection of 50  $\mu$ g of ferric iron, where nigral TH-immunopositive neurons' cell number is dramatically reduced (up to 95 % cell loss; Gerlach and Riederer 1996).

In humans, at least 70 % of dopaminergic neurons in the SN and 70 % of striatal dopamine must be lost before the symptoms of PD develop (Riederer and Wuketich 1976; Halliday et al. 1996). Progressive neurodegeneration thus occurs several years prior to the onset of the characteristic motor symptoms. The current lesion mimics that which occurs in the parkinsonian brain, where nigral cell loss exhibits a distinct pattern, beginning with the loss of the ventrolateral neurons and progressing to loss of dorsolateral and medial nigral cell groups (Halliday et al. 1996). A localized loss of 50 % of ventrolateral dopaminergic neurons thus represents the pattern of cell loss occurring in preclinical PD. The success of this approach is demonstrated by the absence of behavioral changes following apomorphine challenge or a striatal dopamine deficit in these animals (Double et al. 2003).

A nearly 50 % reduction of TH-positive neuronal cells was also found in the SN of adult male Wistar rats 1 week following unilateral intranigral injection of  $2-\mu$ l human NM (0.5 mg/ml; Zecca et al. 2008). This injection also induced a strong inflammatory response assessed by immunostaining with glial fibrillary acidic protein and Iba-1 in close vicinity to the NM granules. It has been recently shown that human NM in vitro induces microglial proliferation, chemotaxis and activates proinflammatory microglial properties by stimulating the release of tumor necrosis factor, interleukine-6, and nitric oxide (Wilms et al. 2003).

# 4 Conclusion

In summary, studies using unilateral intranigral administration of ferric iron and NM-bound ferric iron have been shown to result in dopaminergic neurodegeneration with characteristic features of the human disease in the rat (Table 1). Further, these studies provided insights into the pathophysiological mechanisms of redox-active iron underlying dopaminergic neuronal cell death in PD.

The ideal model of PD (Duty and Jenner 2011) would exhibit a high degree of construct validity (that is similar pathogenesis to the disease), of face validity (that is, similarity in symptoms, neurochemistry, and neuropathology to the human condition) as well as predictive validity (that is, the ability to positively identify agents that are clinically effective). Unilateral administration of ferric iron and NM-bound ferric iron into the SN is therefore a valid experimental model of mechanisms, leading to dopaminergic neurodegeneration in PD, and for the development of neuroprotective agents for PD that stop or slow disease progression (Table 1).

	Unilateral intranigral ferric iron injection	Unilateral intranigral human NM-bound ferric iron injection
Construct validity	High (increased SNc ferric iron, oxidative stress, inflammation)	High (increased SNc ferric iron, oxidative stress, inflammation, proteasome inhibition)
Face validity		
Symptoms	Reduction of spontaneous locomotor activity	Absence of motoric disturbances: subclinical model
Neurochemistry	Severe striatal dopamine depletion (up to 95 %)	Moderate striatal dopamine depletion (50 %)
Neuropathology	Nearly complete dopaminergic SNc cell loss	Dopaminergic SNc cell loss
		Moderate dopaminergic SNc cell loss (50 %)
Progression	Intracellular proteinaceous aggregates: No	Not investigated
	Yes	Not investigated
Predictive validity	Yes (development of iron chelators for protection of dopaminergic neurons)	Not investigated

Table 1 Similarities of ferric iron-induced characteristic features to the human condition

NM neuromelanin, SNc substantia nigra pars compacta

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Molecular, Cellular, and Behavioural Effects Produced by Perinatal Asphyxia: Protection by Poly (ADP-Ribose) Polymerase 1 (PARP-1) Inhibition

T. Neira-Peña, P. Espina-Marchant, E. Rojas-Mancilla, D. Esmar, C. Kraus, V. Munoz, R. Perez, B. Rivera, D. Bustamante, J. L. Valdes, M. Hermoso, P. Gebicke-Haerter, P. Morales, and M. Herrera-Marschitz

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#### Abstract

Perinatal asphyxia implies oxygen interruption at birth, leading to death whenever reoxygenation is not promptly reestablished. Reoxygenation triggers a cascade of biochemical events for restoring function at the cost of improper

Department of Psychopharmacology, Central Institute of Mental Health, Mannheim, Germany

The contribution of TN-P and PE-M has been equally relevant.

T. Neira-Peña • P. Espina-Marchant • E. Rojas-Mancilla • D. Esmar • C. Kraus • V. Munoz • R. Perez • B. Rivera • D. Bustamante • J.L. Valdes • M. Hermoso • P. Morales •

M. Herrera-Marschitz (🖂)

Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile e-mail: mh-marschitz@med.uchile.cl

P. Gebicke-Haerter

Millennium Institute BNI-Chile, Program of Molecular and Clinical Pharmacology, ICBM, BNI, Medical Faculty, University of Chile, Santiago, Chile

homeostasis. The effects observed long after perinatal asphyxia have been explained by overexpression of sentinel proteins, such as poly(ADP-ribose) polymerase 1 (PARP-1), competing for NAD<sup>+</sup> during reoxygenation, leading to the idea that sentinel protein inhibition constitutes a suitable therapeutic strategy. Asphyxia also induces transcriptional activation of proinflammatory factors, including NF $\kappa$ B, and its subunit p65, whose translocation to the nucleus was found here, is significantly increased in brain tissue from asphyxia-exposed animals, in tandem with PARP-1 overactivation, suggesting that PARP-1 inhibition downregulates the expression of proinflammatory cytokines. Indeed, TNF- $\alpha$  and IL-1 $\beta$  were found to be increased 8 and 24 h after perinatal asphyxia in mesencephalon and hippocampus of rat neonates.

The possible neuroprotection effect of nicotinamide has been studied in an experimental model of global perinatal asphyxia in rats, inducing the insult by immersing rat fetuses into a water bath for various periods of time. Following asphyxia, the pups are delivered, immediately treated, or given to surrogate dams for nursing, pending further experiments. Systemic administration of nicotinamide was found to rapidly distribute into the brain reaching a steady-state concentration sufficient to inhibit PARP-1 activity for several hours. Nicotinamide prevented several of the long-term consequences elicited by perinatal asphyxia, supporting the idea that it constitutes a lead for exploring compounds with similar or better pharmacological profiles.

#### Keywords

Behavior • Development • Hypoxia • Neonatal • Obstetric complications • Plasticity • Poly(ADP-ribose) polymerase • Rats • Sentinel proteins

List of Abbreviations	
AIF	Apoptosis-inducing factor
AN 1	Nicotinamide-treated, asphyxia-exposed rats
AS	Asphyxia exposed, saline treated
ATP	Adenosine triphosphate
BAD I	Bcl-2-associated death factor
BAX I	Bcl-2-associated X factor
BBB I	Blood–brain barrier
bFGF I	Basic fibroblast growth factor
CNS (	Central nervous system
COX-2	Cyclooxygenase-2
CREB a	cAMP-response element-binding protein
CS (	Caesarean delivered, saline treated
DG I	Dentate gyros
DNMT1 I	DNA(cytosine-5-)-methyl transferase 1

DPQ	3,4-dihydro-5-[4-(1-piperidinyl)butoxy]-1(2H)-		
DD0010	isoquinolinone		
DR2313	2-methyl-3,5,7,8-tetrahydrothiopyranol[4,3-d]pyrimidine-		
	4-one		
E2F	Family of DNA-binding transcription factors		
EPO	Erythropoietin		
ERCC2	Excision repair cross-complementing rodent repair group 2		
ERK	Extracellular signal-regulated kinases		
FOXO	Subclass of forkhead O family of transcription factors		
FR247304	5-Chloro-2-[3-(4-phenyl-3,6-dihydro-1(2H)-pyridinyl) propyl]-4(3H)-quinazoline		
GAPDH	Glyceraldehyde 3-phosphate dehydrogenase		
HDAC	Histone deacetylases		
HIF	Hypoxia-inducible factor		
HRE	Hypoxia-responsive elements		
ICAM-1	Intercellular adhesion molecule-1		
IGF-1	Insulin-like growth factor-1		
IL-1β	Interleukin-1β		
iNOS	Inducible nitric oxide synthase		
IQ	Intelligence quotient		
ΙκΒ	Inhibitor of kappa B protein		
Ku70	Protein encoded by the XRCC1 gene, required for the		
itu / 0	nonhomologous end joining pathway of DNA repair		
LIG3	DNA ligase 3		
LPS	Lipopolysaccharides		
NAD <sup>+</sup>	Nicotinamide adenine dinucleotide		
NADH	Reduced nicotinamide adenine dinucleotide		
ΝΓκΒ	Nuclear factor-KB		
ONO-1924H	<i>N</i> -3-(4-oxo-3,4-dihydrophthalazin-1-yl)phenyl-4-		
	(morpholin-4-yl) butanamide methane sulfonate		
	monohydrate		
Р	Postnatal day		
p300	Histone acetyltransferase p300		
p65	Transcription factor p65		
PAR	Poly(ADP-ribose) polymers		
PARG	Poly(ADP-ribose) glycohydrolase		
PARPs	Poly(ADP-ribose) polymerases		
PARylated PARP	Poly-ADP-ribosylated PARP		
PBS	Phosphate-buffered saline		
PCAF	P300/CBP-associated factor		
PCr	Phosphocreatine		
PHD	Prolyl hydroxylase domain		
PJ34	[ <i>N</i> -(6-oxo-5,6-dihydrophenanthridin-2-yL)- <i>N</i> , <i>N</i> -		
	dimethylacetamide.HCl]		
	,		

POLB, DNA	Polymerase- $\beta$
ROS	Reactive oxygen species
SIRT	Sirtuin
SRY	Sex-determining region Y
Strep-HRP	Streptavidin-horseradish peroxidase
SVZ	Subventricular zone
TH	Tyrosine hydroxylase
TNF	Tumor necrosis factor
TUNEL	Terminal deoxynucleotidyl transferase dUTP nick end
	labelling
VEGF	Vascular endothelial factor
XRCC1	X-ray cross-complementing factor 1

## 1 Introduction

Obstetric complications are important risk factors for several psychiatric and neurological disorders characterized by a delayed clinical onset. Hypoxia is a common factor of obstetric complications, priming brain development by mechanisms not yet established (Basovich 2010).

Delay in starting pulmonary ventilation at birth implies decreased oxygen saturation in blood and decreased oxygen supply to the brain, which depends upon aerobic metabolism for generating energy and for maintaining the respiratory chain and mitochondrial ATPase activity. Hypoxia implies a switch to glycolysis, which for neurons is a poor metabolic alternative, because of low stores of glucose in brain tissue and deficient ATP output by the glycolysis pathway. Furthermore, glycolysis implies production of lactate, which is accumulated in extracellular compartments, causing acidosis, although there is evidence indicating that lactate can provide a significant source of energy to neurons (Wyss et al. 2011). Prolonged hypoxia further involves suppression of gene expression and translation, favoring the activation of genes for sustained survival, such as hypoxia-inducible factor (HIF) and its target genes (Iyer et al. 1998).

The incidence of perinatal asphyxia is still high, despite improvements of perinatal care (2–6/1,000 term births; De Hann et al. 2006), occurring with higher prevalence in developing countries (Lawn et al. 2010). After asphyxia, infants can develop long-term neurological sequelae, their severity depending upon the extent of the insult. Severe asphyxia has been linked to cerebral palsy, mental retardation, and epilepsy, while mild to severe asphyxia has been associated with attention deficits, hyperactivity, and schizophrenia. No strict correlation has yet been demonstrated between the clinical outcome and the severity and/or the extent of the insult, but the outcome and how the infants recover from the insult are related. In a cohort study monitoring approximately 6,000 children for 8 years, it was demonstrated that resuscitated infants, even if asymptomatic for encephalopathy,

showed an increased risk for low IQ score (Odd et al. 2009). At present, there is no therapeutic strategy to significantly prevent the long-term effects produced by perinatal asphyxia, apart from hypothermia, which still is a controversial issue (Robertson et al. 2012).

## 2 Reoxygenation

While the average  $pO_2$  in tissue is approximately 65 mmHg, it can decline below 5 mmHg following severe hypoxia. Reoxygenation is mandatory for restoration of function and survival. However, functional constraints of the cells affected by reduced oxygen may be exacerbated during reoxygenation, implying oxidative stress and production of proinflammatory cytokines, delaying the onset of proper homeostasis and complete recovery (Hagberg et al. 1996; Shalak et al. 2002; Girard et al. 2009). Reoxygenation can even enhance protein synthesis over normal levels (see Fähling 2009), including ribosomal biogenesis, leading to defects in protein synthesis fidelity, a mechanism associated to several neurodegenerative diseases (Lee et al. 2006). A number of proteins has to be increased upon reoxygenation, triggering (i) cell death, (ii) cell repairing (e.g., basic fibroblast growth factor, bFGF; VEGF, CREB-binding protein; ribosomal protein S<sub>19</sub>; casein kinase 1), and (iii) *neuritogenesis* and *synaptogenesis*, required for maturation of neurocircuitries, also implying neurogenesis. Increased postnatal neurogenesis has been observed in subventricular zones (SVZ) and dentate gyrus (DG) of hippocampus in response to hypoxia/ischemia (Scheepens et al. 2003; Bartley et al. 2005).

The leading role in these early events following hypoxia is taken over by HIF-1 $\alpha$  that is both an oxygen sensor and a transcription factor. After heterodimerization with HIF-1 $\beta$ , it binds to hypoxia-responsive elements (HRE) on promoters of genes like VEGF, iNOS, hemoxigenase-1, and EPO or to the oxygen sensors prolyl hydroxylase domain (PHD) proteins 2 and 3. HIF-1 also binds to other transcription factors, such as X-ray cross-complementing factor 1 (XRCC1) (Green et al. 1992), excision repair cross-complementing rodent repair group 2 (ERCC2) (Sung et al. 1993; Chiappe-Gutierrez et al. 1998; Lubec et al. 2002), and poly(ADP-ribose) polymerases (PARPs), mainly PARP-1 (Amé et al. 2004). The coordinated action of the products of these genes is believed to minimize and compensate for the damage induced by hypoxic conditions, acting on genes involved in glycolysis, oxygen transport, cell survival, and apoptosis (see Correia and Moreira 2010).

#### **3** Sentinel Proteins

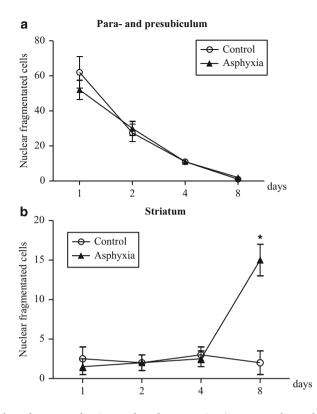
Most of the adaptive changes elicited by hypoxia begin at the level of the DNA. Indeed, within minutes of cerebral hypoxia/ischemia, oxidative damage of DNA is observed in brain parenchyma, including oxidation of guanosine to 8-oxo-guanine and deamination of cytosine to uracil. A major repair mechanism to reconstitute damaged DNA is base excision repair, shown to be elevated following hypoxia/ ischemia, implying activation of PARP-1, XRCC1, ERCC2, DNA polymerase- $\beta$ (POLB), and DNA ligase (LIG3) (see Ellenberger and Tomkinson 2008; Herrera-Marschitz et al. 2011).

PARPs, XRCC1, ERCC2, POLB, and LIG3 are rapidly activated whenever there is a risk of genome damage. These proteins play a pivotal role not only in repairing mechanisms but also in other environmentally induced DNA modifications. They can act as both cell survival- and cell death-inducing factors by regulation of DNA repair, chromatin remodelling, and regulation of transcription. PARP-1 overactivation may be elicited by P300/CBP-associated factor (PCAF), resulting in caspase-independent translocation of apoptosis-inducing factor (AIF) from mitochondria to the nucleus and subsequent cell death by chromatinolysis. Indeed, PARP-1 activation is essential for AIF truncation and release from mitochondria (Kolthur-Seetharam et al. 2006). PARP-1 can be acetylated for enhancing nuclear factor-KB (NFKB)-, FOXO- and Ku70-mediated transcription and for that of tumor suppressor protein p53 (Zampieri et al. 2009). Otherwise, PARP-1 can be deacetylated by proteins of the sirtuin (SIRT) family, histone deacetylases (HDAC) type III, resulting in its inactivation. Activated SIRT also blocks release of truncated AIF from mitochondria (Kolthur-Seetharam et al. 2006; Rajamohan et al. 2009).

PARP-1 activity is strictly controlled to maintain the balance between cell survival and cell death. The equilibrium between the antagonistic actions of SIRT and PARP-1 is maintained through their strict dependency from and competition for NAD<sup>+</sup> (Berger 1985). Increased demand for NAD<sup>+</sup> by PARP-1 inhibits SIRT and vice versa. If both are overactive simultaneously, NAD<sup>+</sup> depletion generates large quantities of nicotinamide that has been shown to inhibit both SIRT and PARP activities (Rajamohan et al. 2009), probably by an end-product inhibitory mechanism. Thus, PARP and SIRT are at the core of epigenetic actions encompassing DNA and histone modifications.

#### 4 Apoptosis

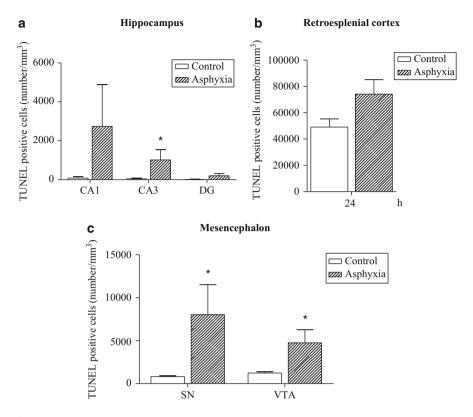
Apoptosis is the prevalent type of delayed cell death in the perinatal brain, mediated by caspase-dependent and caspase-independent mechanisms (Northington et al. 2001; see Morales et al. 2008). Indeed, pro-apoptotic proteins have been observed to be increased following perinatal asphyxia, including Bcl-2-associated X (BAX), and Bcl-2-associated death (BAD) factors, and also anti-apoptotic proteins, including Bcl-2, ERK2, and bFGF, suggesting the activation of neuroprotective and repair pathways (Morales et al. 2008). Extensive and regionally selective nuclear fragmentation has been observed in control and asphycic rat pups, depending upon the stage of development and the analyzed brain region (Dell'Anna et al. 1997). Signs of apoptosis were observed in para- and presubiculum of both control and asphyxia-exposed animals, independently upon the severity of the insult, with the highest number of cells showing nuclear fragmentation observed



**Fig. 1** Number of neurons showing nuclear fragmentation in para- and presubiculum (**a**) and neostriatum (**b**) of asphyxia-exposed and control rat pups 1–8 days after birth. Brain sections (sagittal 20  $\mu$ m serial formalin-fixed sections) were obtained from asphyxia-exposed (20–21 min of asphyxia) (n = 5-7 animals per group) (filled triangles) and the corresponding control (n = 9-16) (open circles) animals, 1–8 days after delivery. The sections were stained for hematoxylin-eosin and in situ DNA double-strand breaks and analyzed under light microscopy at 40× magnification to detect cellular alterations, using Foster atlas (1998) for proper brain region identification (see Dell'Anna et al. 1997). Values are expressed as means ± SEM (\*p < 0.05; Student's *t*-test)

at P1-2, decreasing thereafter, with no cells showing nuclear fragmentation at P8 (Fig. 1a). In contrast, no significant nuclear fragmentation was observed in neostriatum of control pups, but a progressive increase in the number of cells showing chromatin fragmentation was observed in asphyxia-exposed animals, with a maximum observed at P8 (Fig. 1b). Similar results were also observed in frontal cortex and cerebellum, increasing with the length of perinatal asphyxia (Dell'Anna et al. 1997).

The effect of perinatal asphysia on apoptosis is heterogeneous even on a particular region. Indeed, when assayed with an  $Apop Tag^{R}$  TUNEL kit (Millipore, Temecula, CA), the number of apoptotic cells was particularly



**Fig. 2** Delayed cell death following perinatal asphyxia. Twenty-four hours after delivery, asphyxia-exposed and control animals were sacrificed, the brains rapidly removed and fixed in 0.1 M pH 7.4 phosphate-buffered saline (PBS) containing 4 % paraformaldehyde, kept in 20 % sucrose, embedded in cryomatrix (Thermo Electron Corp, Pittsburgh, PA, USA), and stored at  $-80^{\circ}$  C. DNA fragmentation was evaluated in coronal brain sections (20 µm thick) with the TUNEL assay ( $ApopTag^{\circledast}$ ; Millipore, Temecula, CA). TUNEL-positive cells (brown) were manually counted in a Nikon TS100 microscope (magnification  $100 \times$ ). The number of TUNEL-positive cells per mm<sup>3</sup> was determined in sections showing hippocampus (**a**), retrosplenial granular zone (**b**), and mesencephalon (**c**) (Foster atlas 1998) from the asphyxia exposed (*dashed bars*) and the corresponding controls (*open bars*). Comparisons were tested with a Student's *t*-test (\*p < 0.05)

increased in CA1 and CA3 regions of the hippocampus of asphyxia-exposed pups, compared to that from controls (Fig. 2a). A large increase was also observed in the retrosplenial granular cortex (Fig. 2b) (Neira-Peña et al. unpublished), a region connecting the hippocampus with the entorhinal cortex (Wyss and Van Groen 1992). This is an exciting observation, because entorhinal cortex is a brain area consistently observed to be abnormal in psychiatric disorders associated to metabolic insults occurring at birth (Bachus et al. 1997). Indeed, entorhinal cortex (Akil et al. 2000) receives dopamine nerve terminals from mesencephalon

(Hökfelt et al. 1974). In agreement, as shown in Fig. 2c, apoptosis was also increased in mesencephalon following perinatal asphyxia, both in substantia nigra and VTA (Neira-Peña et al. unpublished).

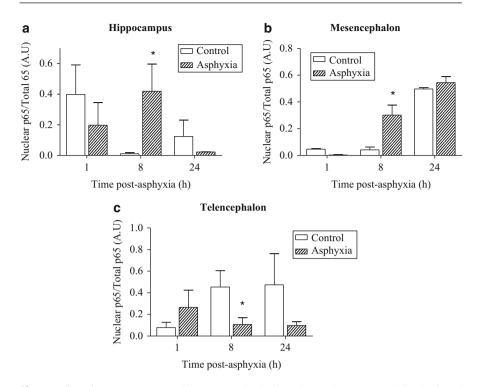
## 5 Neuroinflammation

Asphyxia can induce transcriptional activation of proinflammatory factors, including NF $\kappa$ B that is normally located in the cytoplasm as a heterodimer, composed of p65 and p50 subunits, coupled to I $\kappa$ B. I $\kappa$ B dissociates from the complex when phosphorylated, liberating p65/p50, which are translocated to the nucleus for inducing the transcription of TNF- $\alpha$ , IL-1 $\beta$  intercellular adhesion molecule-1 (ICAM-1), cyclooxygenase-2 (COX-2), iNOS, and IL-6, all genes involved in inflammation (see Girard et al. 2009).

The immature CNS is capable of mounting innate and adaptive immune responses, through microglial and astrocytes, although it has been argued that the immune responses of the developing CNS differ from that of the adult, in part due to immaturity of blood–brain barrier (BBB). Tight junctions are, however, present early during embryonic development (Kniesel et al. 1996), controlling the interchange of proteins, including leukocytes (Engelhardt 2003; Vexler and Yenari 2009). The developing brain is, however, vulnerable to increases of inflammatory cytokines, leading to a substantial cross talk between peripheral and local brain immune components (see Ransohoff et al. 2003; Vexler and Yenari 2009). Indeed, Qiao et al. (2001) have demonstrated that in neonatal brain, disruption of the BBB to proteins occurs earlier after hypoxic-ischemic insult than in the mature brain.

Microglia provide immunosurveillance to the brain by stimulus-dependent activation (see Vexler and Yenari 2009). Microglia populate the developing brain at birth, and activated microglia can release a number of cytokines, including IL-1 $\beta$ , IGF-1, and TNF- $\alpha$ . It is not clear yet whether microglia activation contributes or protects against the long-term deficits induced by perinatal asphyxia, but it has been reported that minocycline, a tetracycline derivative with anti-inflammatory properties, protects the neonatal brain against ischemia, partly by inhibiting microglia activation and monocyte infiltration (Arvin et al. 2002; Dommergues et al. 2003; Denker et al. 2007). Nevertheless, to investigate the involvement of different cytokines is complex, depending upon the type of the insult, upon the timing along the development axis, and also upon specific brain regions.

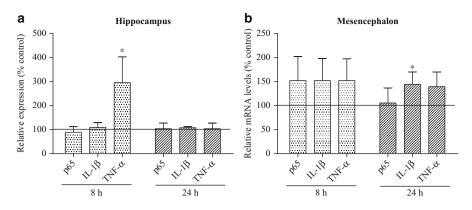
The translocation of NF $\kappa$ B was investigated in tissue samples from mesencephalon, hippocampus, and telencephalon from asphyxia-exposed and control rat pups, 1–24 h after birth, with an antibody specific to p65. It was found (Neira-Peña et al. unpublished) that translocation of p65 to the nucleus was significantly increased (>10×) in mesencephalon of both control and asphyxia-exposed pups 24 h after birth, but that increase was already remarkable 8 h after birth in asphyxia-exposed animals (Fig. 3b). In hippocampus, translocation of p65 was enhanced 1 h after birth in control animals, decreasing thereafter at 8–24 h (Fig. 3a). In contrast, in asphyxia-exposed animals, translocation of p65 was remarkably enhanced in



**Fig. 3** *Effect of reoxygenation on p65 activation.* Asphyxia and control rats were sacrificed 1, 8, and 24 h post-birth; the brain was rapidly removed, taking up tissue samples from hippocampus (**a**), mesencephalon (**b**), and telencephalon (**c**), placed in Eppendorf tubes, frozen in liquid nitrogen, and stored at -80 °C, pending further analysis. Cytosolic and nuclear protein was extracted with a ProteoJet<sup>TM</sup> kit (Fermentas). P65 levels were determined in cytosolic and nuclear protein fractions of tissue samples from asphyxia-exposed (*dashed bars*) and controls (*open bars*) rat pups. Western blot was performed using an anti-NFkB p65 antibody (65 KDa) (Santa Cruz Biotechnology, Inc, Santa Cruz, CA). Tubulin and histone H4 were used as controls for the respective fractions. A density-graphic analysis of the bands was performed. Data are represented as nuclear p65/total p65 ratio of Arbitrary Units (AU) (comparisons were analyzed with a Student's *t*-test; \**p* < 0.05)

hippocampus 8 h after birth, decreasing thereafter to almost negligible levels at 24 h (Fig. 3a). In telencephalon, translocation of p65 increased (>10×) from 1 to 8–24 h after birth in the controls, but in asphyxia-exposed animals, translocation of p65 was increased compared to the controls at 1 h, decreasing thereafter (Fig. 3c). Furthermore, TNF- $\alpha$  mRNA levels were increased in hippocampus 8 h after perinatal asphyxia (Fig. 4a), while IL-1 $\beta$  mRNA levels were only increased in mesencephalon, 24 h after the insult (Fig. 4b).

DNA damage and cell death elicited by perinatal asphyxia has the potential of activating PARP-1, promoting NF $\kappa$ B activation (see Skaper 2003; Gagne et al. 2008). In agreement, Ullrich et al. (2001) showed that the microglial migration towards the site of neuronal injury is controlled by PARP-1 overactivation, correlating with NF $\kappa$ B translocation. To evaluate the role of



**Fig. 4** *mRNA expression of inflammatory-related genes.* Total RNA was extracted with DNAzol, treated with DNAse, and subsequently verified for optical density 260/280 absorption ratios. The integrity of RNA was analyzed by denaturing gel electrophoresis. Reverse transcription was performed with oligo-dT primers from total RNA to obtain cDNA. Relative mRNA levels to GAPDH of p65, IL-1 $\beta$ , and TNF- $\alpha$  were measured by RT-qPCR in tissue samples from hippocampus (**a**) and mesencephalon (**b**). Figure shows the ratio asphyxia-exposed/control rats, at 8 (*stippled bars*) and 24 h (*dashed bars*) after birth. Results, in triplicates, were analyzed with a MxPro software (comparisons tested by a Student's *t*-test; \*p < 0.05)

PARP-1 in inflammatory processes, PARP-1  $^{-/-}$  mice were challenged with lipopolysaccharides (LPS), finding that the knockout mice were resistant to the endotoxin shock compared with wild-type animals. The knockout mice also showed low levels of TNF- $\alpha$ , due to reduced transcriptional activity of NF $\kappa$ B (Petrilli et al. 2004).

Inhibition of PARP-1 downregulates the expression of proinflammatory cytokines and suppresses NF $\kappa$ B-dependent gene transcription in microglia. Kauppinen et al. (2006, 2009) demonstrated in a model of bilateral carotid occlusionreperfusion in rats that treatment with the PARP-1 inhibitor PJ34 ([*N*-(6-oxo-5, 6-dihydrophenanthridin-2-yl)-*N*, *N*-dimethylacetamide.HCl]), 48 h later, rapidly suppressed the ischemia-induced microglial activation, enhancing long-term neuronal survival, neurogenesis, and spatial memory. Also, in a model of forebrain ischemia, PJ34 produced a near-complete inhibition of microglia activation and a significant reduction of neuronal death in the hippocampus (Hamby et al. 2007). Furthermore, it was reported that minocycline protects cortical neuron cultures against genotoxic agents causing DNA damage. Interestingly, minocycline also inhibits PARP-1 activity (Alano et al. 2006).

#### 6 Epigenetics

PARP-1 activity is under control of and controls in a reciprocal way the activity of H1-histone, resulting in exclusion of H1 from a subset of PARP-regulated promoters and subsequent regulation of chromatin structure.

PARP itself can be poly-ADP-ribosylated (PARylated PARP), which is reversed by poly-ADP-ribose glycohydrolase (PARG). There is a link between poly-ADPribosylation and DNA methylation implying DNA silencing (Cohen-Armon et al. 2007). Specifically, it has been shown that PARG overexpression, or reduction of PARylated PARP, results in methylation of CpG islands in the DNA (cytosine-5-)methyltransferase 1 (DNMT1) promoter (Zampieri et al. 2009). Gene silencing leads to widespread DNA hypomethylation, resulting in increased gene expression. Conversely, when PARylated PARP occupies the DNMT1 promoter, it protects its unmethylated state and DNMT transcription. Additionally, other transcription factors, such as p300, SRY, or E2F, are associated with PARP at the DNMT1 promoter (Rajamohan et al. 2009).

Thus, epigenetics is a factor to be considered when studying plastic changes elicited by metabolic insults occurring at birth. There is evidence that the molecular machinery that regulates histone acetylation and DNA methylation might be intimately involved in synaptic plasticity, including learning and memory. Importantly, dysfunction of epigenetic gene expression in the brain may result in neurodegenerative and psychiatric diseases (Sananbenesi and Fischer 2009). Conversely, drugs that increase histone acetylation (such as HDAC inhibitors) exhibit neuroprotective and neuro-regenerative properties in animal models of neurodegenerative diseases, enhancing cognitive functions (MacDonald and Roskams 2009).

# 7 A Therapeutic Target

The adverse consequences of long-term effects of overactive PARP-1 in response to asphyxia (Martin et al. 2005) have led to the notion that PARP-1 is a suitable target for therapeutic interventions preventing the long-term effects of perinatal asphyxia (see Herrera-Marschitz et al. 2011). Nicotinamide has been proposed as a prototype for counteracting PARP-1 overactivation (see Virag and Szabo 2002), replacing NADH/NAD<sup>+</sup> (Zhang et al. 1995), and protecting against oxidative stress (Yan et al. 1999; Wan et al. 1999; Sakakibara et al. 2000) and inflammation (Ducrocq et al. 2000).

It has been reported (Bustamante et al. 2003, 2007) that nicotinamide prevents several of the changes induced by perinatal asphyxia on monoamine contents and dopamine release monitored with in vivo microdialysis 3 months after birth, even if the treatment was delayed for 24 h, suggesting a clinically relevant therapeutic window, supporting the idea that nicotinamide can constitute a therapeutic strategy against the long-term deleterious consequences of perinatal asphyxia, as already proposed for several other pathophysiological conditions (see Virag and Szabo 2002). Nicotinamide has also been proposed as a treatment of Alzheimer's disease (Green et al. 2008).

The therapeutic doses of nicotinamide (0.8 mmol/kg, i.p.) used in the abovereported studies produce a long-lasting inhibition of PARP-1 activity measured in brain and heart from asphyxia-exposed and control animals (Allende-Castro et al. 2012) and also a decrease of the number of apoptotic nuclei in hippocampus, increased by perinatal asphyxia (Morales et al. 2010).

The use of nicotinamide has, however, been challenged because of its low potency, limited cell uptake, and short cell viability, stimulating the search for more specific compounds, such as 3-aminobenzamide (Ducrocq et al. 2000; Hortobagyi et al. 2003; Koh et al. 2004); 3,4-dihydro-5-[4-(1-piperidinyl)butoxy]-1(2H)-isoquinolinone (DPQ) (Takahashi et al. 1999); PJ34 (Abdelkarim et al. 2001); *N*-3-(4-oxo-3,4-dihydrophthalazin-1-yl)phenyl-4-(morpholin-4-yl) butanamide methane sulfonate monohydrate (ONO-1924H) (Kamanaka et al. 2004); 5-chloro-2-[3-(4-phenyl-3,6-dihydro-1(2H)-pyridinyl) propyl]-4(3H)-quinazoline (FR247304) (Iwashita et al. 2004); and 2-methyl-3,5,7,8-tetrahydrothiopyranol[4,3-d]pyrimidine-4-one (DR2313) (Nakajima et al. 2005). Ultrapotent novel PARP inhibitors are in clinical trials for reducing parenchymal cell necrosis following stroke and/or myocardial infarction, downregulating multiple pathways of inflammation and tissue injury following circulatory shock, colitis, or diabetic complications (Jagtap and Szabo 2005).

There is, however, concern about applying ultrapotent PARP inhibitors during development, since it has been shown that PARP is required for efficient repair of damaged DNA (Trucco et al. 1998; Schultz et al. 2003), suggesting that moderate PARP-1 inhibitors should be chosen for neuronal protection, whenever used for pediatric patients (Moonen et al. 2005; Geraets et al. 2006).

Nicotinamide is therefore an interesting molecule because of its low potency, which can be an advantage when used for developing animals, antagonizing the effects elicited by PARP-1 overactivation without impairing DNA repair or cell proliferation. Nicotinamide has already been tested in human clinical trials without showing any significant toxicity, although its therapeutic efficacy is still controversial (Macleod et al. 2004). Nicotinamide can constitute a lead for exploring compounds with similar or better pharmacological profiles.

## 8 An Experimental Model for Perinatal Asphyxia

A model for investigating the short- and long-term outcome of perinatal asphyxia was proposed at the Karolinska Institutet, Stockholm, Sweden, in the nineties (Bjelke et al. 1991; Andersson et al. 1992; Herrera-Marschitz et al. 1993). Asphyxia is induced at the time when the rats are delivered providing some features with clinical relevance: (i) it occurs at term, (ii) it is largely noninvasive, (iii) it allows studying peripheral and brain tissue in the same animal, (iv) it allows studying short- and long-term consequences of the insult in the same preparation, and (v) it is highly reproducible among laboratories. Lubec and coworkers in Vienna (Austria) (Lubec et al. 1997a, b; Seidl et al. 2000) have stressed the issue that the model allows to study the early phase of perinatal asphyxia, as observed in the clinical setup.

The model starts by evaluating the estral cycle of young female Wistar rats  $(\sim 2 \text{ months of age})$ , in order to plan for a programmed mating. A vaginal frottis is taken for evaluating the cycle, identifying proestrus, estrus, metaestrus, or diestrus. The female is then exposed to a male at the time of the proestrus for one night, evaluating thereafter for the presence of a vaginal clot. Thus, the time of delivery is calculated, supported by ethological and clinical observations, to predict the exact time of delivery (22 days). At the time of delivery, a first spontaneous birth can be observed before the dams are neck dislocated, and subjected to a caesarean section and hysterectomy. The uterine horns containing the fetuses are immediately immersed into a water bath at 37° C for various periods of time (0–22 min). Following asphyxia, the pups are removed from the uterine horns and resuscitated by cleaning from fluid and amniotic tissue, freeing the mouth and the nose. Pups exposed to caesarean delivery only (CS, 0 asphyxia) or to mild asphyxia (2–10 min) are rapidly resuscitated, without requiring anything else but removing fluid and amniotic tissue from the mouth. For pups exposed to longer periods of asphyxia (19-21 min), resuscitation implies expert and skilful handling, and it takes a long time (4–6 min) for a first gasping and even longer time for establishing a more or less regular breathing, always supported by gasping. After 60 min of caretaking, the pups are given to surrogate dams for nursing, pending further experiments. An Apgar scale is applied during the recovery period, similar to that observed in neonatal units, including parameters such as weight, sex, color of the skin, respiratory frequency, gasping, vocalization, muscular rigidity, and spontaneous movements (Dell'Anna et al. 1997; Morales et al. 2010; see Herrera-Marschitz et al. 2011). The Apgar evaluation is a critical parameter, because it assesses whether the pups are subjected to mild or to severe asphyxia, which is directly determined by the percentage of survival and recovery (Herrera-Marschitz et al. 1993, 1994). The Apgar evaluation also provides information about the condition shown by the caesarean-delivered control pups, which has to be similar to that shown by vaginally delivered pups. Thus, the Apgar evaluation is a requirement when using the present model of perinatal asphyxia, because it permits to compare results obtained by different laboratories and/or different treatments. The quality of the handling of the pups and the experience of the surrogate dam are important factors for the acceptance and nursing of both asphyxia-exposed and control pups.

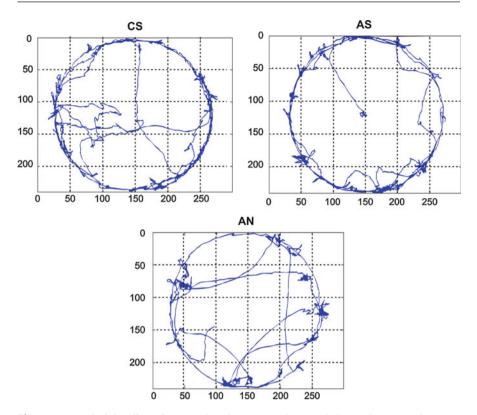
The model has been useful for describing early molecular, metabolic, and physiological effects. Tissue sampling can be collected immediately after delivery, the time when the pups are removed from the uterine horns (0 min, with or without previous immersion into a water bath), or soon after reoxygenation (Lubec et al. 1997a, b; Seidl et al. 2000; Engidawork et al. 2001). Thus, energy-rich phosphates have been measured in brain and peripheral tissue immediately after delivery following short or long periods of perinatal asphyxia, demonstrating that adenosine triphosphate (ATP)/phosphocreatine (PCr) levels were first significantly decreased in kidneys (after 2 min of asphyxia), then in brain tissue (after 10 min), but in heart ATP dropped down after 20 min of asphyxia (Lubec et al. 1997a; Seidl et al. 2000; see Herrera-Marschitz et al. 2011).

## 9 Effect of Nicotinamide on the Long-Term Functional Consequences Elicited by Perinatal Asphyxia

Motor and cognitive alterations of variable severity, including cerebral palsy, seizures, spasticity, attention deficit, hyperactivity, mental retardation, and/or neuropsychiatric syndromes with delayed clinical onset have been associated to perinatal asphyxia (du Plessis and Volpe 2002; Van Erp et al. 2002; Kaufman et al. 2003; Vannuci and Hagberg 2004; Odd et al. 2009). With the proposed experimental model, several labs have investigated the behavioral effects associated to perinatal asphyxia, addressing motor function (Bjelke et al. 1991; Chen et al. 1995), emotion (Dell'Anna et al. 1991; Hoeger et al. 2000; Venerosi et al. 2004, 2006; Simola et al. 2008; Morales et al. 2010), and spatial memory (Boksa et al. 1995; Iuvone et al. 1996; Hoeger et al. 2000, 2006; Loidl et al. 2000; Van de Berg et al. 2003; Venerosi et al. 2004).

The issue whether perinatal asphyxia produces long-term effects on cognition has also been investigated. Novel object recognition was studied with a test applied under ethological-like conditions, devoid of any stressful cues or primary reinforcers, such as swimming, food deprivation, or electric shocks (Ennaceur and Delacour 1988). The animal is tested to discriminate between objects differing in shape and color, without any genuine significance to the rat or being previously associated to any rewarding or aversive stimuli. During a first session, two copies of the same object are presented to the rat for 4 min and then again, during a second session, when one of the previously presented objects is replaced by a novel one, similar in size, but different in shape and/or color. The rat has to recognize the novel object, spending longer time exploring the novel than that previously presented. The first and the second sessions are separated by different time intervals, for evaluating learning consolidation. A good memory would be able to recognize a previously presented object after a long time elapsing between a first and a second session, meaning that the animal would concentrate on exploring the novel object. Novel object recognition was studied using a 15 or 60 min interval, when the animals were 3 months old (Simola et al. 2008). No differences were observed between asphyxia-exposed (20 min asphyxia) and control animals when a 15 min interval elapsed between the first and the second session. Both asphyxia-exposed and control rats recognized the novel stimulus similarly well, spending longer time exploring the novel object. However, when 60 min elapsed between the first and second session, asphyxia-exposed animals spent less time exploring the novel object, indicating that asphyxia-exposed rats could not recognize its novelty. Conversely, asphyxia-exposed rats could not remember that one of the objects was already presented during the first session (Simola et al. 2008). This was a straightforward experiment showing a subtle consequence of hypoxia occurring at birth, impairing a cognitive function that shows up only after a proper challenge. It is very much reminiscent to the clinical experience revealing effects only when the child starts the primary school (see Odd et al. 2009; Strackx et al. 2010).

In a parallel study, the same condition was repeated, but a cohort of asphyxiaexposed and of control rats received 0.8 mmol/kg, i.p. of nicotinamide 1, 24, and 48 h after birth and evaluated 2–3 month later for the novel object recognition



**Fig. 5** *Reversal of the effect of perinatal asphysia on exploratory behavior by neonatal nicotinamide treatment.* Exploratory behavior was examined in a circular arean placed in a noise-free room. Arean was 140 cm in diameter painted in black, with a continuous wall of 15 cm high and placed 50 cm above the floor. A camera placed 250 cm over the arean was used for recording the rat performance. The camera was connected to a computerized processing unit (*CPU*) placed in the next room where the examiner stayed during the experiment. At 2–3 months of age, control (CS) and asphysia-exposed (AS) rats treated with saline (3 × 0.1 ml/kg, i.p) or with nicotinamide (AN) (0.8 mmol/kg, i.p.; 1, 24 and 48 h after birth) were placed in the center of the arena free to explore for 5 min (Individual experiments taken from cohorts reported by Allende-Castro et al. 2012)

task. Nicotinamide did not affect the performance of the control animals, but prevented the effect of perinatal asphyxia, since the animals receiving nicotinamide performed identically as the controls on novel object recognition (Morales et al. 2010).

Asphyxia-exposed, treated, and control animals were tested in the first 2–3 months after birth for exploratory behavior, monitored with a video camera when animals were placed in a circular arena in a noise-free room. Each animal was left to explore the new environment for 5 min, monitoring the time expended in the center of the arena, exploiting the natural aversion of rodents for open spaces devoid of thigmotactic cues. A decreased time in the center of the arena was considered as a measure of anxiety (Perez de la Mora et al. 2012). Figure 5 shows the tracking of

a caesarean-delivered saline-treated (CS), a saline-treated asphyxia-exposed (AS), and a nicotine-treated asphyxia-exposed (AN) rat. As previously reported (Allende-Castro et al. 2012), the AS, but not CS or AN, rats avoided areas devoid of thigmotactic cues, indicating unconditioned fear and anxiety, also observed when analyzing the behavioral profile on an elevated plus maze of parallel experimental cohorts (Morales et al. 2010). Nicotinamide prevented the anxiety-related behavior in both experimental paradigms.

## 10 Pharmacodynamics and Pharmacokinetics of Nicotinamide

It was further investigated whether the effect of nicotinamide was related to PARP-1 inhibition using an enzymatic assay of PARP-1 activity, based on the accumulation of PAR polymers. It was found that a single dose of 0.8 mmol/kg, i.p. of nicotinamide decreased PARP-1 activity by  $\sim$ 70 % in brain (mesencephalon, telencephalon) and peripheral (heart) tissue of asphyxia-exposed and caesarean-delivered control pups, compared with the corresponding saline controls, 1-24 h after birth. A remarkable PARP-1 inhibition was still observed 24 h after the treatment. In vivo microdialysis experiments in  $\sim$ 8h-old pups allowed to monitor the distribution of a single dose of nicotinamide (0.8 mmol/kg, i.p.), in peripheral (subcutaneous) and brain (neostriatum) tissue (Allende-Castro et al. 2012), finding a rapid distribution of nicotinamide into the brain, detectable for longer than 6 h at a steady-state concentration of 20  $\mu$ M. In the same study, theophylline was also investigated, only finding PARP-1 inhibition in peripheral tissue, in agreement with a poor distribution of a single dose of the ophylline (0.14 nmol/kg, i.p.) into the brain compartment.

#### 11 Conclusion

Perinatal asphyxia is still a health concern worldwide, a risk factor for several mental and neurological disorders with a delayed clinical onset. Hypoxia implies a severe energetic crisis, leading to death if reoxygenation is not promptly restored. The functional constraints produced by the lack of oxygen can be exacerbated by and during the reoxygenation period, implying oxidative stress, synthesis, and release of metabolic by-products delaying the onset of proper homeostasis and recovery. A leading role is played by HIF-1 $\alpha$ , promoting the synthesis of proteins though to minimize or compensate the damage induced by the hypoxic condition, but also to promote proinflammatory cytokines leading to cell death. A number of sentinel proteins are rapidly activated whenever there is a risk of genome damage, stimulating base excision repair. PARP-1 has been shown to play a pivotal role for repairing damaged DNA, but also for eliciting caspase-independent cell death when repairing is not viable, modulating pro- and anti-inflammatory signalling.

Furthermore, there is equilibrium between PARP-1 and SIRT proteins, regulating histone acetylation and DNA methylation, the core of epigenetic modification.

PARP-1 overactivation can lead to NAD<sup>+</sup> exhaustion, worsening the energy crisis, leading to the hypothesis that PARP-1 is a suitable target for therapeutic interventions preventing the long-term effects of perinatal asphyxia, nicotinamide being a prototype for counteracting PARP-1 overactivation.

The neuroprotection effect of nicotinamide has been studied in an experimental model of global perinatal asphyxia in rats. In this model, asphyxia is induced by immersing rat fetuses into a water bath for various periods of time. Following asphyxia, the pups are delivered, immediately treated, or given to surrogate dams for nursing, pending further experiments.

Following systemic administration, nicotinamide rapidly distributes into the brain and peripheral compartments reaching a steady-state concentration sufficient to inhibit PARP-1 activity for several hours. Nicotinamide prevents several of the long-term consequences elicited by perinatal asphyxia, supporting the idea that it can constitute a lead for exploring compounds with similar or better pharmacological profiles.

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# MPTP: Advances from an *Evergreen* Neurotoxin

# Anna R. Carta, Augusta Pisanu, Carlos Barcia, and M. T. Herrero

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A.R. Carta (🖂)

Department of Biomedical Sciences, University of Cagliari, Cagliari, Italy e-mail: acarta@unica.it

A. Pisanu CNR Institute of Neuroscience, Cagliari, Italy e-mail: augustapisanu@tiscali.it

C. Barcia

Clinical and Experimental Neuroscience / Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (NiCE–CIBERNED), School of Medicine, University of Murcia, Murcia, Spain e-mail: barcia@um.es

#### M.T. Herrero

Clinical and Experimental Neuroscience (NiCE–CIBERNED), School of Medicine, Campus de Espinardo, University of Murcia, Murcia, Spain

Centro de Investigación Biomédica en Red de Enfermedades Neurodegenerativas (CIBERNED), School of Medicine, University of Murcia, Murcia, Spain

Predepartmental Unit of Medicine, School of Health Sciences, Universitat Jaume I, Castellón, Spain e-mail: ezquerro@med.uji.es; mtherrer@um.es

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#### Abstract

Since its discovery in 1976, 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) models in rodents and nonhuman primates have continuously renewed to keep up with progresses of Parkinson's disease (PD) research. MPTP is able to reproduce almost all the clinical and neuropathological features of PD when administered to monkeys. In contrast, up to date no rodent model has been able to reproduce all PD features in one. Nevertheless, MPTP is a very versatile neurotoxin that can reproduce different aspects of PD pathology, depending upon the dose and regimen of administration. At the present time, a number of different MPTP models have been developed, allowing researchers to investigate either the classical PD neuropathology and neuroprotective mechanisms or known pathological processes underlining more recently discovered aspects of the disease, such as nonmotor symptoms. In this chapter primate and rodent MPTP models are reviewed, focusing mainly on the contribution that different MPTP protocols can offer to reproduce the multifaceted aspects of the disease.

#### Keywords

Model • MPTP • Parkinson • Primate • Rodent

## 1 Introduction

Preclinical investigation of Parkinson's disease (PD) neuropathology and testing of the neuroprotective strategies depends on the availability of reliable experimental animal models. However, modeling PD represents a sorely intricate task. Although several neurotoxin-based and genetic models of PD have been developed, most of them display limitations. Optimally, experimental models of PD should display face validity, reproducing the highest number of features of human PD; predictive validity is also sought, where the neuropathology should evolve progressively as in PD, and therapy responses should be comparable to the human disease; lastly, on a neuropathological view, a PD model should display construct validity, reproducing the complex interaction between neurons and surrounding cells. 1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) is able to reproduce almost all the clinical and neuropathological features of PD when administered to monkeys (Wichmann and DeLong 2003). However, up to date no rodent model has been able to reproduce all of these features together. Depending on the

MPTP model, important limits may include lack of neurodegeneration, absence of motor symptoms despite a nigrostriatal degeneration, failure to reproduce the progressive, and slow nature of human pathology. Moreover, PD involves multiple brainstem nuclei and systemic pathologies, which is difficult to reproduce in a single animal model. On the other hand, MPTP has conveyed to be a very versatile neurotoxin, reproducing different aspects of PD pathology, depending upon the dose and regimen of administration adopted. Keeping this concept in mind and carefully pre-considering dosing regimens, MPTP models provide an incredible contribution to the understanding of the neuropathological mechanisms of PD and to the investigation of neuroprotective and disease-modifying strategies.

# 2 PD Symptoms and Neuropathology

Parkinson's disease is the most frequent neurodegenerative disorder after Alzheimer disease, affecting nearly 1 % of the population over 60s. Symptoms are mainly motor, including akinesia, rigidity, bradykinesia, postural abnormalities, and tremor. Furthermore, over the past decades substantial evidence has documented a plethora of nonmotor aspects of the disease. These may have either a central or an autonomic basis and include olfactory dysfunction, REM behavior disorders, chronic constipation, orthostatic intolerance, sweating, swallowing difficulties, and chronic fatigue. In addition, neuropsychiatric components such as depression, anxiety, cognitive decline, and dementia may often complicate the clinical picture. Most motor symptoms of PD are caused by the degeneration of nigrostriatal dopamine neurons. In addition, the involvement of brain areas other than the dopaminergic has been described since many years and is now gaining increasing attention for the understanding of biochemical mechanisms of nonmotor symptoms. Hence, PD is now recognized as a complex multisystem pathology, where neurodegeneration extends to brain areas such as the locus coeruleus, raphe nuclei, and dorsal motor nucleus of the vagus as well as to parts of the autonomic nervous system (Hawkes et al. 2009; Braak and Del Tredici 2008).

The appearance of motor symptoms reflects the nigrostriatal degeneration, rising when striatal DA drops to 20–30 % of physiological levels and when over 60 % of nigral neurons are lost. Most nonmotor symptoms also correlate with the advancing disease; however, some of them such as sleep behavior disorder and olfactory deficits precede the onset of motor symptoms by a number of years, being currently investigated as markers for early clinical diagnosis (Chaudhuri and Naidu 2008).

Alongside the degeneration of neurons of the substantia nigra *pars compacta* (SNc), cellular hallmarks of PD are the Lewy bodies and dystrophic neurites. Moreover, a number of malfunctioning cellular processes have been described, including oxidative stress, mitochondrial dysfunction, and impaired protein disposal system (Olanow 2007). In the past decade, several evidences have pointed to neuroinflammation as a key pathological feature and likely a hallmark of PD (Long-Smith et al. 2009). Postmortem studies have described reactive microglia around degenerating neurons in the SN, while inflammatory cytokines levels are elevated in the brain, cerebrospinal fluid, and serum of PD patients

(Blum-Degen et al. 1995; Dobbs et al. 1999; McGeer and McGeer 2008; Mogi et al. 1994). In PD, activated microglia are engaged in a vicious cycle called reactive gliosis, in which constant inflammation contributes to neuronal damage (Carta et al. 2011; Hirsch and Hunot 2009; Ransohoff and Cardona 2010; Schwartz et al. 2006).

## 3 MPTP Neurotoxicity

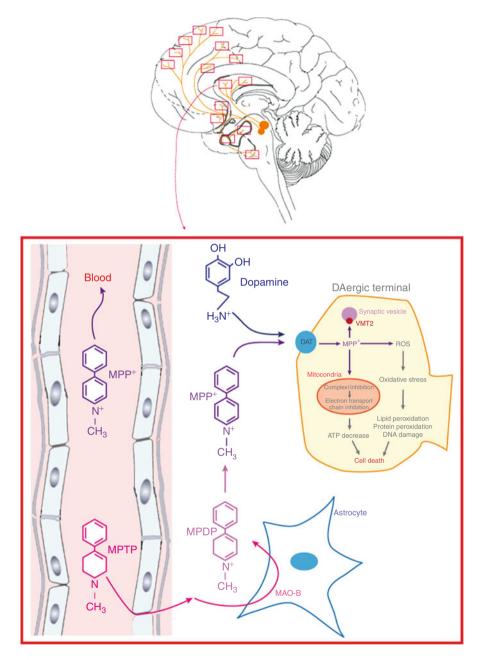
The use of MPTP to model PD is one of the few cases where the effect of a neurotoxin was discovered in humans first and then translated to an animal model. In 1982 a cluster of drug addicts in California were inadvertently intoxicated by MPTP, found as a by-product in a street preparation of an analog of the opioid meperidine. The parkinsonian syndrome shown was almost indistinguishable from PD, characterized by severe bradykinesia and rigidity, which were relieved by traditional antiparkinsonian therapies (Langston et al. 1983).

MPTP does not exert the neurotoxic effect by itself, but, after crossing the blood-brain barrier by virtue of its lipophilic nature, is rapidly oxidized to 1-methyl-4-phenyl-2,3dihydropyridinium (MPDP+) by monoamine oxidase B (MAO-B) in glia and serotonergic neurons (Fig. 1). It is then converted by spontaneous oxidation to the neurotoxic compound 1-methyl-4-phenylpyridinium (MPP+), which is a high-affinity substrate for the dopamine transporter (DAT). MPP+ can now effortlessly be taken up into dopaminergic cells and concentrated within mitochondria, following the mitochondrial transmembrane potential (Javitch et al. 1985; Mayer et al. 1986; Ramsay and Singer 1986; Ransom et al. 1987) (Fig. 1). At this level, MPP+ exerts its toxic effect by inhibiting the complex I of the electron transport chain and impairing oxidative phosphorylation (Nicklas et al. 1985). This event induces a decrease in ATP (Chan et al. 1991; Fabre et al. 1999) content and increase in ROS production that rapidly leads to cell damage and cell death (Hasegawa et al. 1990). Besides these direct effects on dopaminergic neurons, an important contribution to MPP+ toxicity is given by glial cells, which not only are responsible of MPP+ production but also are activated by MPP+ itself or by injury factors released by degenerating neurons, to upregulate inflammatory cytokines and the inducible nitric oxide synthase (iNOS) (Teismann et al. 2003; Youdim et al. 2002). Nitric oxide (NO) easily crosses cell membranes and diffuses toward surrounding dopaminergic neurons, thus contributing to oxidative stress, since peroxynitrite (ONOO-) derived from the reaction of NO with superoxide is highly reactive and induces nitration of proteins (Ischiropoulos and al-Mehdi 1995; Przedborski et al. 1996, Przedborski and Vila 2003).

## 4 Animal Species and Strand Sensitivity to MPTP Neurotoxicity

## 4.1 Rodents

In general, rodents are less susceptible than primates to MPTP toxicity. Among rodents, mice are mostly used, while rats are not recommended for



**Fig. 1** Once MPTP has crossed the BBB, it is transformed to MPP+ inside the astrocytes through the action of MAO-B. Then, MPP+ enters DA neurons through the DAT and either inhibits the complex I of the mitochondria, inducing cell death, or enters into the synaptic vesicles through the VMAT-2. The levels of astrocytic MAO-B and the expression of DAT and VMAT-2 in DA neurons are important factors in the susceptibility to MPTP

MPTP models, as they display an elevated resistance to the neurotoxin (Giovanni et al. 1994; Heikkila et al. 1984; Sahgal et al. 1984). Variable susceptibility among mouse strains has also been found, the C57black/6 strain being the most sensitive to systemic MPTP injection and currently the most popular for PD research (Giovanni et al. 1994; Schmidt and Ferger 2001; Sedelis et al. 2000). Such differences among animal species and strains may be partially related to the metabolic processes that convert MPTP to toxic metabolites in the brain and periphery. MPTP clearance in the brain, and liver metabolism to nontoxic species, is slower in black mice as opposed to white mice and rats (Riachi et al. 1988). Furthermore, MAO activity in brain microvessels is lower in mice, leading to a reduced conversion of MPTP to MPP+ before entering the brain and to an enhanced capacity of MPTP to reach its sites of toxicity (Riachi et al. 1989). A higher VMAT2 density in DA neurons of rats with respect to mice has been also found (Staal and Sonsalla 2000). Based on that, a more efficient sequestration into synaptic vesicles has been suggested as a further mechanism of protection and reduced vulnerability of rats against MPP+ neurotoxicity. More recently, it has been suggested that variances in immunological backgrounds, or differences in glial and microglial cell number and ratio, might be major contributing factors in susceptibility differences to MPTP (Smeyne et al. 2005; Yasuda et al. 2008).

## 4.2 Primates

Apparently, all species of primates seem to be susceptible to the neurotoxin MPTP; however, species and primate families display some differences. For instance, old-world primates (those evolved in Africa, Asia, and small areas in Europe) such as baboons and macaques show a higher degree of susceptibility as compared to new-world primates (those evolved in America), such as capuchins monkeys, squirrel monkeys, or marmosets (Emborg 2007).

In order to achieve a stable parkinsonism in old-world primates such as baboons, macaques, green monkeys, and vervet monkeys, doses of MPTP range from 0.2–0.3 mg/kg to 0.6–1 mg/kg. However, 1.2 mg/kg, 2 mg/kg, or even higher doses should be used for new-world primates (Barcia et al. 2004a; Elsworth et al. 2000; Emborg 2007; Hantraye et al. 1996; Herrero et al. 1993b; Pearce et al. 1995; Perez-Otano et al. 1991; Varastet et al. 1994). These differences have led to believe that the higher degree in the phylogenetic scale, the more susceptibility to MPTP is shown. Hence, humans are highly susceptible to MPTP according to the small doses extrapolated from accidentally intoxicated individuals (Langston et al. 1983). We do not have data from great apes, such as gorillas, orangutans, or chimpanzees to compare with, but specific characteristics, described below, suggest that they may also present high susceptibility.

Different factors may contribute to the different susceptibility among primate species. Specifically, the levels of MAO-B and neuromelanin seem to play a main role. According to its higher susceptibility, primate brain contains high proportions of MAO-B (about 70–80 %) as compared with rodent brain that contains

predominantly MAO-A (Zesiewicz and Hauser 2002). On the other hand, neuromelanin, a pigment present in dopaminergic neurons, is contained in higher levels in the mesencephalic dopaminergic areas most susceptible to MPTP (Herrero et al. 1993b). In addition, elegant experiments have shown that chloroquine, a drug that blocks MPP+ binding to neuromelanin, protects monkeys from MPTP-induced dopaminergic degeneration (D'Amato et al. 1987). It is well known that MPTP susceptibility increases with age. Besides the aging effect on DA neurons itself, neuromelanin accumulation with age may also increase neuronal vulnerability to MPTP (Herrero et al. 1993a). It is noteworthy that within primates, there is a progressive increase of neuromelanin pigmentation as the phylogenetic relationship to man becomes closer (Marsden 1961; Scherer 1939), in line with an increased susceptibility to MPTP of phylogenetically higher primate species.

## 5 Currently Available In Vivo MPTP Models

Although a number of MPTP paradigms involving variable routes of administration have been developed to date, a main limit for all of these models is the failure of reproducing the complete and multifaceted PD pathology, particularly when working with rodents. For this reason, none of them are eligible as the best preclinical PD model, but each one should be regarded and carefully chosen for the specific purpose of the study, e.g., investigating motor or nonmotor disturbances, neurodegenerative and neuroprotective mechanisms, and neuroplastic changes.

## 5.1 Rodents

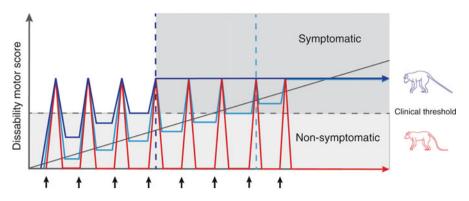
A variety of models are currently available, mostly in mice, since their higher sensitivity to systemic MPTP, including systemic administration (i.e., subcutaneous, intraperitoneal, or intranasal), or local injection (i.e., intraventricular or intranigral). Subcutaneous or intraperitoneal MPTP administration is used in mice only, while intracerebral injection or intranasal administration, allowing MPTP to bypass the BBB, is used in both mice and rats (Prediger et al. 2010; Yazdani et al. 2006). With regard to the protocol of administration, rodent models can be referred to as acute, subacute, or chronic. Acute models include single day-treatment paradigms. MPTP hydrochloride is administered via peripheral route and may involve a single dose of 20-40 mg/kg or a series of four doses (10-30 mg/kg) usually injected in 2 h intervals (Bezard et al. 1997a; Jackson-Lewis et al. 1995; Kohutnicka et al. 1998). A recent acute MPTP model involves the intranasal neurotoxin administration, where mice or rats receive small amounts (10–15  $\mu$ l) of a MPTP solution directly in each nasal cavity (Prediger et al. 2006, 2010; Rojo et al. 2006). Subacute paradigms generally last about a week, with a daily administration protocol of doses ranging from 20 to 30 mg/kg (Carta et al. 2009; Tatton and Kish 1997; Vila et al. 2000). Chronic MPTP paradigms might be more reliable PD models, since they are able to mimic a higher number of PD features, including the slow progressive pattern of neurodegeneration seen in human PD. Chronic protocols can last 3–5 weeks, with continuous or intermittent MPTP treatment. Intermittent treatment involves either daily or twice/week neurotoxin peripheral injection at doses 4–20 mg/kg (Bezard et al. 1997a; Lau et al. 1990; Petroske et al. 2001; Schintu et al. 2009). Concomitant administration of the clearance inhibitor probenecid (MPTPp model), by reducing brain and kidney excretion of MPTP, helps in keeping higher and more stable levels of the neurotoxin during treatment, resulting in a slowly progressing but persistent neurodegeneration (Lau et al. 1990; Petroske et al. 2001; Schintu et al. 2009). An alternative route of chronic neurotoxin administration consists in the continuous MPTP delivery via osmotic minipumps providing a daily dose of 30 mg/kg (Fornai et al. 2005).

## 5.2 Primates

The most common model used in primates to induce a parkinsonian syndrome is the intravenous administration of MPTP through the saphenous vein (Emborg 2007; Guridi et al. 1996). With this type of administration, a bilateral and stable parkinsonism can be achieved. Another commonly used protocol is the intracarotid administration, consisting of MPTP infusion through the carotid artery, leading to a hemi-parkinsonism, where animals only show symptoms of PD in one side of the body (contralateral to the injection side) (Emborg 2007; Kordower et al. 2000). The majority of animals display symptoms after 3–4 doses of MPTP, but some individuals may present a clear parkinsonian syndrome with even one dose (highly susceptible animals), while others may not show signs even after several doses (not susceptible animals) (Fig. 2). For this reason, it is important to evaluate and monitor each animal individually.

Both intrasaphenous and intracarotideal models show advantages as well as disadvantages. Parkinson's symptoms may be better measured in hemi-parkinsonian monkeys since the evaluation can be compared with the contralateral side of the same animal. On the other hand, the bilateral model, although trickier to detect small changes, mimics closer the clinical scenario observed in humans. However, in the bilateral model, one side is generally slightly more affected than the other, making it difficult to assess regular motor scores for parkinsonian monkeys. As a main advantage, the bilateral model with intrasaphenous infusion does not involve any surgery, being therefore a less complex and safer procedure for the animal.

The protocols of MPTP administration can be considered acute, subacute, subchronic, and chronic. Acute treatment is usually referred to a single dose of MPTP, with animals being sacrificed within 24 h, although some variations can be found in the literature. Subacute models may be more variable, being described as a short-term MPTP administration, with animals being sacrificed after few weeks (Garrido-Gil et al. 2009; Vazquez-Claverie et al. 2009a). A subchronic protocol is commonly referred to a MPTP treatment lasting few weeks to a month and sacrificing the animals within 3 months (Barcia et al. 2004a; Goldstein et al. 2003). The chronic protocol, also called remote, is usually referred to



**Fig. 2** The panel on the right shows the typical posture of a control longtail macaque (Macaca fascicularis) compared with a parkinsonian macaque. Note that the parkinsonian monkey shows a particular curvature of the limbs and trunk as well as rigidity of the tail. The panel on the left shows a chart with the typical curve of the motor score of a susceptible monkey compared with an asymptomatic one. After the systemic effects, the susceptible monkeys maintain stable parkinsonian symptoms, while asymptomatic animals recover completely after the systemic effects

a MPTP treatment lasting 3 months or more and sacrificing the monkeys within 6–35 months or even years after MPTP administration (Barcia et al. 2004b; Colosimo et al. 1992; Goldstein et al. 2003; Halliday et al. 2009; McGeer et al. 2003).

## 6 Motor and Nonmotor Symptoms

MPTP administration induces a short-term and transient systemic effect both in monkeys and in rodents, which is independent from its ability to alter locomotor activity but may be a confusing factor for the detection of PD symptoms. Systemic effect may include piloerection, hypersalivation, hypokinesia and seizures, tachy-cardia, hyperventilation, and hypothermia that may require the use of a heater and recover within 24–48 h. Hours after MPTP administration, primates may also show some episode of vomiting and lack of appetite that may result in loss of body weight. When working with MPTP, it is essential to respect the time of quarantine to reach a full elimination of any residual MPP+ by the urine and feces.

### 6.1 Rodents

The induction and detection of motor impairments in mice are profoundly affected by the schedule of MPTP delivery and behavioral tests applied, for which reason results have been variable and sometimes ambiguous (Table 1). Acute MPTP paradigms convey a reduction, no changes, or even an increase in locomotor activity (Colotla et al. 1990; Fredriksson and Archer 1994; Luchtman et al. 2009;

MPTP protocol	Motor activity		Nonmotor symptoms	
Acute	Colotla et al. 1990 ↑		Anderson et al. 2007 (enteric)	
	Fredriksson and Archer 1994	↓	Monaghan et al. 2010 (cognitive)	
	Rousselet et al. 2003	=	Prediger et al. 2010 (olfactory, cognitive)	
			Tian et al. 2008 (enteric)	
Subacute	Arai et al. 1990	↓	Laloux et al. 2008 (sleep)	
	Chia et al. 1996	Î		
	Petroske et al. 2001	-		
	Willis and Donnan 1987	-		
Chronic	Bezard et al. 1997b	=	Schintu et al. 2009 (olfactory)	
	Fornai et al. 2005	↓	-	
	Petroske et al. 2001	Ļ		
	Rousselet et al. 2003	$\uparrow$		
	Schintu et al. 2009	Ļ		

Quinn et al. 2007; Rousselet et al. 2003; Sedelis et al. 2000; Tillerson et al. 2002). In any case, the motor impairment induced by acute MPTP recovers nearly completely after 3–7 days postinjection (Schmidt and Ferger 2001). Subacute MPTP treatments have also given contradictory results. When mice are treated for a period ranging from 5 days to few weeks, no changes (Petroske et al. 2001; Willis and Donnan 1987), decrease (Arai et al. 1990), or increase (Chia et al. 1996; Luchtman et al. 2009) in locomotor activity has been reported.

After a series of improvements and protocol adjustments, chronic MPTP models are now able to provide more consistent results. In a first attempt of reproducing the disease in chronic nature, MPTP was given daily at a very low dose for 20 days (Bezard et al. 1997a). Unfortunately, despite a progressive decline of TH immunoreactivity, the MPTP-treated mice did not display parkinsonian motor abnormalities after this dosing regimen (Bezard et al. 1997b). In the last decade, the chronic MPTPp protocol, firstly proposed by Lau in 1990 and widely characterized by several studies thereafter, brought a significant improvement to PD models (Lau et al. 1990). First, the MPTPp administration schedule (twice/week) allows to perform motor tests between injections, offering the useful advantage of monitoring motor symptoms in the presence of an ongoing neurodegeneration. When tested along the treatment, mice display an increasing impairment in motor performance, suggesting the development of progressive nigrostriatal degeneration (Schintu et al. 2009). Moreover, MPTPp-treated mice display a reduced motor performance when tested 3 weeks to 6 months after treatment, suggesting persistent neurodegeneration (Petroske et al. 2001; Schintu et al. 2009). Confirming dopaminergic degeneration, amphetamine-induced locomotor activity is reduced after 4 weeks from neurotoxin treatment (Al-Jarrah et al. 2007). As an alternative model, the continuous MPTP delivery through minipumps was reported to dramatically depress spontaneous activity in the open field test (Fornai et al. 2005). However, constant brain levels of MPTP obtained by this route may induce some nonspecific systemic effects, including depressed motor behavior that may confound the assessment of PD symptoms.

A further burst to the use of MPTP models in PD research comes from recent studies, highlighting that specific protocols may be suitable for studying nonmotor aspects of the disease (Table 1). The possibility of mimicking symptoms associated with the early premotor phases of PD has been deeply explored in the intranasal MPTP infusion model (Prediger et al. 2006, 2010). In this model, animals develop selective symptoms such as olfactory dysfunction and cognitive deficits, while not displaying a clear motor impairment, suggesting that this MPTP protocol may provide a useful model of premotor symptoms of PD. Interestingly, in the MPTPp model, olfactory deficits appear after the first neurotoxin administration, preceding motor symptoms, accordingly with the graded progression of PD neuropathology (Hawkes et al. 2009; Haehner et al. 2011; Schintu et al. 2009). Cognitive impairment was reported after an acute intraperitoneal MPTP administration as well (Monaghan et al. 2010). Among other nonmotor symptoms, sleep alterations have been reported after a subacute MPTP treatment (Laloux et al. 2008). Peripheral pathologies associated with PD have been sparely investigated. Interestingly, behavioral and electrophysiological defects in the enteric nervous system, associated with a significant reduction of TH-positive neurons in the same site, were reported after acute MPTP intoxication in mice (Anderson et al. 2007; Tian et al. 2008). Certainly more efforts are sought to evaluate the possible contribution of MPTP models to the understanding of peripheral aspects of PD pathology; nevertheless, these few studies support the interesting possibility that MPTP may provide a useful tool to investigate on the graded neuropathology hypothesis of the disease (Hawkes et al. 2009).

## 6.2 Primates

The evidence of major motor symptoms of PD after MPTP intoxication of young drug addicts prompted researchers to test the neurotoxin in nonhuman primates. Since then, the MPTP model has been used both as a pathogenic and as a clinical symptomatic model in monkeys for its peculiarity of mimicking almost all of the clinical hallmarks of the disease (Bezard and Przedborski 2011), including the side effects induced by dopamine replacement with levodopa and/or dopaminergic agonists (Blanchet et al. 2004; Herrero et al. 1995; Jenner 2003). After MPTP intoxication, monkeys develop akinesia/bradykinesia, rigidity, and freezing phenomena. They also may show tremor in the arms and hands and occasionally in the head. Differently from humans, who show resting tremor, MPTP-intoxicated monkeys show a tremor that appears with action. Eventually, monkeys may display occasional falls and vertical saccadic eye movements. After MPTP exposure, a stable parkinsonism may be observable through the years (Halliday et al. 2009). Although some animal may improve slightly during the observation, parkinsonism

is usually irreversible. However, as described above, it is important to consider that some individuals are not susceptible to MPTP, and after the recovery of systemic symptoms (24–48 h), no apparent motor alterations may be seen. MPTP-treated monkeys also show a postural alteration that is characterized by the curvature of the limbs and trunk and a peculiar rigidity of the tail (Fig. 2).

MPTP-treated monkeys also show similarities with PD patients with regard to nonmotor symptoms. Similarly to the patients, MPTP-treated monkeys present alterations in the heart sympathetic innervation (Goldstein et al. 2003; Li et al. 2002; Yuste et al. 2012), enteric nervous system (Chaumette et al. 2009), retina (Cuenca et al. 2005), as well as sleep disorders (Almirall et al. 1999) and modifications in the circadian rhythm of secretion of hormones such as prolactin and melatonin (Barcia et al. 2003). Moreover, when the schedule of MPTP intoxication is chronic and slow, fronto-executive cognitive impairments are evident even before and independent of the motor symptoms (Schneider and Pope-Coleman 1995). The first cognitive symptoms include significant impairment in performing short- and medium-duration delay trials, suggesting attentional deficits that disrupt the encoding of behaviorally relevant information to be kept in short-term memory. These symptoms have been also reported in asymptomatic MPTP-treated monkeys. Moreover, after repeated administration of low doses of MPTP for long periods, parkinsonian monkeys develop significant impairment in planning, cognitive flexibility, and attention set shifting (Decamp and Schneider 2004). Altogether, these evidences point to the MPTP primate model as a useful tool for studying the progression of parkinsonian symptoms (Storvik et al. 2010) as well as for the understanding of the broad range of systems that are affected in PD (Olanow et al. 2011).

Importantly, the MPTP primate model faithfully reproduces L-DOPA-induced dyskinesia. Similarly to humans, L-DOPA administration in parkinsonian primates brings about recovery of motor symptoms, while the prolonged L-DOPA treatment causes the appearance of dyskinesia. The MPTP primate model gives the unique opportunity to replicate a dyskinesia nearly identical to parkinsonian patients (Ahmed et al. 2010; Clarke et al. 1987; Herrero et al. 1995).

# 7 Neuropathological Markers of PD in MPTP Models

While nigrostriatal degeneration can be achieved by different protocols of MPTP administration, the dosing regimen and schedule of administration are the main determinants for modeling histological features of PD, such as  $\alpha$ -synuclein-positive inclusion bodies, chronic neuroinflammation, and neuronal death by apoptotic mechanisms.

## 7.1 Neurodegeneration in Rodents

Classically, MPTP leads to the selective degeneration of nigrostriatal dopamine neurons. Striatal dopamine terminals are extremely sensitive to MPTP toxicity, and

**Table 2** Effect of different MPTP protocols on striatal dopamine levels (DA), TH immunoreactivity (TH-IR), histological staining, and the appearance of apoptotic cells in the substantia nigra pars compacta (SNc).  $\downarrow$  indicates a decrease in the measured parameter, = indicates no changes with respect to drug-naïve mice, - means not analyzed

MPTP protocol		DA (Str)	TH-IR (SNc)	Histological staining (SNc)	Apoptotic nuclei (SNc)
Acute	Jackson-Lewis et al. 1995	-	Ļ	Ļ	No
	Hallman et al. 1985	Ļ	=	_	_
	Heikkila et al. 1984	Ļ	$\downarrow$	_	_
	Ricaurte et al. 1986	Ļ	=	=	_
Subacute	Carta et al. 2009	_	Ļ	$\downarrow$	_
	Petroske et al. 2001	Ļ	=	=	_
	Tatton and Kish. 1997	_	$\downarrow$	$\downarrow$	Yes
	Vila et al. 2000	_	Ļ	_	Yes
	Willis and Donnan 1987	Ļ	=	-	_
Chronic	Alvarez-Fischer et al. 2008	$\downarrow$	Ļ	_	_
	Bezard et al. 1997a, b	_	Ļ	_	_
	Fornai et al. 2005	Ļ	Ļ	_	_
	Novikova et al. 2006	Ļ	$\downarrow$	_	Yes
	Petroske et al. 2001	Ļ	Ļ	Ļ	_
	Rousselet et al. 2003	Ļ	$\downarrow$	_	_
	Schintu et al. 2009	$\downarrow$	Ļ	$\downarrow$	_

even a single dose leads to a rapid decrease of striatal dopamine and DOPAC levels (Hallman et al. 1985; Jackson-Lewis et al. 1995; Schintu et al. 2009) (Table 2). Moreover, an important aspect that has been revaluated in recent years is the MPTP-induced depletion of catecholamines in areas other than the striatum. MPTP-induced decrease of dopamine and/or noradrenaline in the olfactory bulb and prefrontal cortex, associated with the appearance of typical PD nonmotor symptoms, supports the interesting perspectives of using MPTP to model nonmotor aspects of the disease (Dluzen 1992; Dluzen and Kefalas 1996; Hallman et al. 1985; Luchtman et al. 2009; Prediger et al. 2010; Tanila et al. 1998).

As compared to striatal terminals, degeneration of cell bodies in the SNc is a more challenging task. MPTP toxicity toward dopaminergic cell bodies is age dependent, particularly after acute protocols (Ricaurte et al. 1987). Moreover, contradictory results have been reported on nigral lesion after acute (Hallman et al. 1985; Heikkila et al. 1984; Jackson-Lewis et al. 1995; Ricaurte et al. 1986), subacute (Carta et al. 2009; Petroske et al. 2001; Vila et al. 2000; Willis and Donnan 1987), and chronic (Alvarez-Fischer et al. 2008; Bezard et al. 1997a; Fornai et al. 2005; Petroske et al. 2001; Rousselet et al. 2003; Schintu et al. 2009) MPTP regimens (Table 2). To this regard, the possible lack of correlation between the loss of TH-positive immunoreactivity and loss of Nissl-stained cells is a pivotal factor to be considered (Bian et al. 2007; Jackson – Lewis et al. 1995; Seniuk et al. 1990). For instance, after acute MPTP treatment, the number of TH-positive neurons decreases rapidly, while the effective loss of Nissl-stained cells only begins at 12 h postinjection and continues up to 4 days (Jackson-Lewis et al. 1995). Therefore, neurodegeneration is best shown through histological direct evidence. Accordingly, a lack of correlation between motor impairment and loss of nigral TH-positive neurons has been often reported. Importantly, the rate and mechanism of cell death is also strictly dependent on the MPTP schedule and dosing (Table 2). After acute and subacute protocols, cell death occurs rapidly. In contrast, chronic MPTP, particularly when associated with a clearance inhibitor, induces a slower and progressive loss of DA neurons (Alvarez-Fischer et al. 2008; Petroske et al. 2001; Schintu et al. 2009). Furthermore, neuronal loss is a transient phenomenon after short-term regimens, while on the other hand, it is long lasting after chronic regimens, being stable up to 6 months after the last MPTP administration (Petroske et al. 2001; Schintu et al. 2009). Finally, acute MPTP protocols result in a rapid non-apoptotic, likely necrotic cell death, while subacute and chronic regimens, where lower MPTP doses are delivered over days or weeks, lead to preferential cell death by apoptotic mechanisms (Jackson-Lewis et al. 1995; Novikova et al. 2006; Tatton and Kish 1997; Vila et al. 2000).

## 7.2 Neurodegeneration in Primates

The degeneration of the SNc in MPTP-treated monkeys presents a regiondependent vulnerability, similarly to humans. The ventrolateral part is the most susceptible, displaying the highest degree of dopaminergic degeneration (80–90 %). This area is equivalent to the  $\alpha + \beta$  area defined in humans, presenting very similar features, and it is equally vulnerable in humans and in all families of primates. Other mesencephalic areas are less susceptible and appear to be partially protected against degeneration. The ventral tegmental area (VTA), the dorsal and lateral part of the SNc, presents a 50 % of dopaminergic loss upon MPTP intoxication, while the periaqueductal gray matter (PAG) and periretrorubral area A8 shows 10 % or less dopaminergic degeneration (Herrero et al. 1993b). These differences in regional susceptibility may be due to several factors such as the level of neuromelanin, as described above, as well as the levels of calbindin, DAT, and vesicular monoamine transporter-2 (VMAT-2) that also show regional differences within dopaminergic areas. Accordingly, dopamine neurons of most susceptible mesencephalic areas display high levels of DAT and low levels of VMAT-2, showing the highest cytoplasmic accumulation of MPP+ (Dauer and Przedborski 2003).

The loss of DA neurons of the SNc in MPTP-treated monkeys mimics and matches with the scenario observed in human PD. Usually, there is a fine correlation between the level of dopaminergic cell loss caused by MPTP and motor alterations (Barcia et al. 2011; Mounayar et al. 2007). However, this correlation may have a range of variation depending on the adaptation, recovery, and compensatory mechanisms that the animal may develop in response to MPTP.

In contrast with DA neurons, the degeneration of dopaminergic fibers in the striatum seems to differ between humans and nonhuman primates. In PD patients, the putamen is the most affected striatal area. However, MPTP seems to affect to a higher extent the caudate rather than the putamen in primates (Moratalla et al. 1992; Snow et al. 2000). Moreover, other brain regions may be affected by MPTP in primates as, for instance, the locus coeruleus (Forno et al. 1986; Herrero et al. 1993a) or the dorsal raphe nucleus (Perez-Otaño et al. 1991). It has also been reported that the damage and alteration of Purkinje cells of the cerebellum may correlate with nigral degeneration (Heman et al. 2012; Necchi et al. 2004). Although the cerebellar activity is altered in PD patients, histological evidences in this nucleus are elusive.

## 7.3 Alpha-Synuclein/Lewy Body Deposits in Rodents

Ubiquitin- and  $\alpha$ -synuclein-positive Lewy bodies have not been observed with any MPTP treatment in mice, despite the evidence that  $\alpha$ -synuclein is involved in MPTP neurotoxicity in this species (Dauer et al. 2002; Shimoji et al. 2005). Nevertheless, an increase of  $\alpha$ -synuclein-positive neurons in the SNc of mice treated with a subacute MPTP regimen has been reported (Vila et al. 2000). Moreover, in mice chronically treated with MPTP either i.p. or through minipumps, alpha-synuclein-immunoreactive inclusions have been reported in a few neurons of the SNc by two studies (Fornai et al. 2005; Meredith et al. 2002) but not by other studies (Alvarez-Fischer et al. 2008; Shimoji et al. 2005), leaving this issue controversial.

## 7.4 Alpha-Synuclein/Lewy Body Deposits in Primates

 $\alpha$ -Synuclein aggregation after the administration of MPTP occurs in both old-world (Kowall et al. 2000) and new-world primates (Purisai et al. 2005). This phenomenon has been related with the dopaminergic cell death, although the mechanism linking  $\alpha$ -synuclein aggregation with neuronal damage is unclear (Sulzer 2010). Despite  $\alpha$ -synuclein accumulation, studies performed in acute MPTP-treated primates have revealed that this does not result in the formation of Lewy bodies (Kowall et al. 2000; Purisai et al. 2005). Since these experiments have been performed in acute protocols of MPTP administration, it has been postulated that a longer period of years may be required for Lewy body formation. However, studies performed in monkeys bearing parkinsonism from over a decade failed to reveal any Lewy body formation (Halliday et al. 2009), which suggests that this phenomenon may be an exclusive feature of the human disease.

## 7.5 Neuroinflammation in Rodents

A number of studies have reported a neuroinflammatory response in mice treated either acutely (Kurkowska-Jastrzebska et al. 1999), subacutely (Carta et al. 2009),

or chronically (Schintu et al. 2009) with MPTP. However, the mouse MPTP model presents some limitations in reproducing the inflammatory scenario that characterize PD. In vivo and ex vivo observations in parkinsonian patients, as well as in MPTP-treated primates, have shown a chronic and persistent inflammatory response, where activated microglia is engaged in the auto-fueling and detrimental cycle named reactive gliosis (Hirsch and Hunot 2009). In contrast, the neuroinflammatory response in mice is transient and tightly related to the neuro-toxin administration (Schintu et al. 2009).

Nevertheless, MPTP studies in mice have contributed to highlight the involvement of neuroinflammation in dopaminergic degeneration. Upon MPTP administration, neuroinflammation has been mainly described as microglia activation associated with the overproduction of inflammatory cytokines. Acute and subacute MPTP intoxication causes a rapid and transient microglial activation in the striatum and in the SNc that returns to basal levels within 14 days post-intoxication (Carta et al. 2009; Kurkowska-Jastrzebska et al. 1999). Moreover, a transient increase in either protein or mRNA for inflammatory cytokines in both areas was reported shortly after MPTP injection (Ciesielska et al. 2003; Kurkowska-Jastrzebska et al. 1999; Lofrumento et al. 2011; Pattarini et al. 2007). The chronic administration of MPTP in association with probenecid induces a gradual increase of microglial activation and density (Meredith et al. 2005; Novikova et al. 2006; Schintu et al. 2009). Interestingly, activated microglia persist longer after chronic MPTP intoxication as compared to shorter protocols, being still detectable up to 6 months posttreatment (Meredith et al. 2005; Schintu et al. 2009). In addition, microglia undergoe changes in morphology and phenotype upon this protocol of MPTP administration, gradually switching to more activated states, associated with altered cytokines production (Pisanu et al. in preparation).

## 7.6 Neuroinflammation in Primates

Different from mice, the inflammatory response caused by MPTP in primates is persistent and may be perpetuated over the years (Barcia et al. 2004b; McGeer et al. 2003). This inflammatory response is characterized by an increase of proinflammatory cytokines in the blood and brain, similar to PD patients, and can be clearly detected at the cellular level (Barcia et al. 2005, 2011; Hunot and Hirsch 2003). From a histological point of view, the inflammatory response is characterized by microglia and astroglia activation specifically in the SNc, appearing particularly higher in the  $\alpha + \beta$  area where DA cell loss is higher (Barcia et al. 2004b). After MPTP intoxication, microglial cells display a bigger cell body and a higher number of processes and ramifications and overexpress human leukocyte antigen-DR (HLA-DR) in their membrane (Barcia et al. 2009; McGeer et al. 2003). These features are very similar or even identical to the scenario observed in humans intoxicated with MPTP and in PD patients (Langston et al. 1999; McGeer et al. 1988). Importantly, the level of activation appears very similar in acute and chronic protocols of MPTP administration (Vazquez-Claverie et al. 2009b). On the other

hand, astrocytes also appear over-reactive in MPTP-treated primates similarly to humans, showing a bigger cell body and increased number of ramifications (Barcia et al. 2009; Forno et al. 1992). Particularly, the glial fibrillary acidic protein (GFAP) is overexpressed in reactive astrocytes over the dopaminergic pathway, and again, it is particularly increased in the  $\alpha + \beta$  area of the SNc, where DA loss is highest. Importantly, the levels of cytokines in the blood and brain seem to contribute to the inflammatory response (Barcia et al. 2011; De Lella Ezcurra et al. 2010; Pott Godoy et al. 2008). Particularly, IFN- $\gamma$  and TNF- $\alpha$  are thought to be involved in the mechanisms of microglial and astroglial activation and may be key cytokines in the neuroinflammatory response occurring in dopaminergic degeneration (Barcia et al. 2011; De Lella Ezcurra et al. 2010; Mount et al. 2007).

## 8 Conclusion

MPTP administration to primates reproduces almost all of the clinical hallmarks of PD, being used as a pathogenic as well as clinical symptomatic model of the disease. Importantly, MPTP-induced parkinsonism in monkeys is usually irreversible, and the neurodegeneration progresses even after withdrawal of neurotoxin. On the other hand, to reproduce PD by MPTP administration in rodents represents a more intricate task. Up to date, no MPTP rodent model has been able to reproduce all PD features in one. Nevertheless, a number of protocols have been developed that can reproduce different aspects of PD pathology, depending upon the dose and regimen of MPTP administration. Therefore, while rodent models allow to investigate on motor and nonmotor symptoms and on PD neuropathology, a careful evaluation of the most appropriate protocol for the study purpose has to be made in advance. Keeping this concept in mind, MPTP models of PD have provided and continue to give a great contribution to the understanding of disease neuropathology and to the development of new symptomatic and disease-modifying compounds.

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# Nanomedicines for Nervous System Diseases

# Dongwei Guo, JoEllyn M. McMillan, and Howard E. Gendelman

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Re	References					

D. Guo

e-mail: dongwe.guo@unmc.edu

J.M. McMillan • H.E. Gendelman (🖂)

e-mail: jmmcmillan@unmc.edu; hegendel@unmc.edu

Departments of Pharmaceutical Sciences, University of Nebraska Medical Center, Omaha, NE, USA

Department of Pharmacology and Experimental Neuroscience, University of Nebraska Medical Center, NE, USA

#### Abstract

The need for the development of nanomedicines for the treatment of degenerative and inflammatory diseases of the nervous system is of immediate importance. Nanoparticles are drug carriers that can specifically target diseased brain subregions and thus enhance therapeutic efficacy. Multifunctional nanoparticles delivering pharmaceutical agents serve a dual purpose: While they facilitate drug delivery they also provide imaging of cell dynamics and pharmacokinetic assessments. Polymer science and nanotechnology could be used to improve diagnostic and therapeutic outcomes. Despite their promise in combating disease, nanomedicines have yet to make their mark as broadly viable pharmaceuticals. The translation of nanoformulation technology for clinical therapeutic and diagnostic applications shows both real potential and inherent obstacles. However, safety concerns remain operative for their potential broad clinical implications, tempering any translational potential.

#### **Keywords**

Cell-based drug delivery • Nanomedicines • Nanoparticles • Neuroimmunology • Neuroinflammatory disease

## 1 Introduction

A range of insults can affect the nervous system that include infection, trauma, vascular insufficiency, misfolded aggregated protein-induced degeneration, toxins and metabolic dysregulations (de Vries et al. 1997, 2012; Frank-Cannon et al. 2009; Graves and Vernino 2012; Kanwar et al. 2012a; Rosenberg 1997). All lead to activation of the immune system that serves as a key factor in both the etiology and progressive nature of a number of neurological diseases (de Vries et al. 1997, 2012; Frank-Cannon et al. 2009; Rosenberg 1997). Such immune responses can serve as disease propagators and healers. The former leads to sustained phenotype switches for glia (microglia and astrocytes) with the secretion of a broad range of inflammatory mediators that include excitotoxins, pro- and anti-inflammatory cytokines, and chemokines (Ransohoff and Engelhardt 2012). These result in alterations in phagocytosis, intracellular killing, clearance, and antigen presentation. Ultimately compromise of the blood-brain barrier (BBB) ensues with extravasation of toxic proteins, altered glial secretions, and neurotoxicity (de Vries et al. 1997; Kanwar et al. 2012a; O'Callaghan et al. 2008; Rosenberg 1997). While complex in nature, each component step for disease provides opportunities for therapeutic intervention (Kanwar et al. 2012b). In all, the targeting of such interventional strategies with facilitation of drug penetrance across the BBB could lead to improved therapeutic outcomes in central nervous system (CNS) disease (Kanwar et al. 2012a).

## 1.1 Current Treatment Strategies

Treatment of Alzheimer's and Parkinson's diseases (AD and PD, respectively), amyotrophic lateral sclerosis (ALS), a range of microbial infections and multiple sclerosis (MS) is limited to ameliorating symptoms and as such poses continuous challenges to clinicians and clinical investigators (de Vries et al. 1997, 2012; Kanwar et al. 2012a). Simply stated, conventional treatments are palliative. Curative therapies are available, for the most part, only for infectious diseases (Elman et al. 2007; Nowacek et al. 2009a). Even with the potential of disease cure, microbial resistance remains an ever significant concern. Furthermore, accurate diagnosis often precludes early therapeutic intervention. Complicating treatment for neurodegenerative, neuroinfectious, and neuroinflammatory diseases is poor penetration of many drugs across the BBB. To this end, considerable efforts are being placed on developing novel strategies for improving drug penetrance and delivery to sites within the brain parenchyma that are disease targets as well as developing improved methods for early disease diagnosis (Alam et al. 2010).

## 1.2 Urgent Need for Nanomedicines

Considerable emphasis is being placed on the future development of therapeutic drug formulations that target the CNS and halt ongoing toxic, metabolic, and inflammatory events that affect the tempo and progression of neurodegenerative diseases (Graves and Vernino 2012; Ulbrich and Lamprecht 2010). Nanotechnology offers one important means toward such a goal. The ability of nanoparticles to traverse the BBB can be exploited to enhance drug entry into the CNS (Fernandes et al. 2010; Kanwar et al. 2012b). Nanoparticles targeted to the CNS can be efficient delivery strategies to improve the efficacy of commonly used medicines along with developing new therapeutics intended to reach the site of disease action (Kanwar et al. 2012b; Metcalfe and Fahmy 2012; Silva 2005; Ulbrich and Lamprecht 2010). In addition, nanoparticle-based imaging systems can provide earlier and more effective diagnosis of subtle brain dysfunctions (Kanwar et al. 2012b; Stoll and Bendszus 2010).

## 2 Nanoparticle Compositions

Disorders of the CNS that include AD, PD, ALS, and stroke are rapidly increasing in disease prevalence and incidence (Brown et al. 2005; Checkoway et al. 2011; Mayeux 2003). Therapies that can change the course of disease would have substantial impact as AD alone costs more than \$100 billion/year in the United States for medical and institutional care. Such numbers will certainly double in the next decades (ADI 2012). Currently, disease diagnosis is made, for the most part, by clinical observation as it has for more than 100 years (de Carvalho et al. 2008;

Jack et al. 2011; Li et al. 1991; Rao et al. 2006; Yew and Cheng 2009). Drugs, vaccines, or regenerative proteins present "real" possibilities to positively alter disease outcomes.

Delivery of drugs to treat CNS diseases poses particular challenges. Barriers to CNS drug delivery include CNS structure, function, and physiology. To enter into the brain parenchyma, drugs must cross the endothelial cell lining of the microvasculature (Nowacek et al. 2009a). Under physiologic conditions, this barrier selectively regulates intracellular and paracellular exchange of nutrients, macromolecules, and cells between the blood and the CNS. Permeability of many drugs through the BBB is very low even though the capillary surface area is large. Thus, the BBB plays an indispensable role in regulating drug CNS concentrations (Alam et al. 2010).

Development of ways to overcome the barriers to CNS drug delivery is of clinical relevance. Delivery of drugs by nanocarrier systems is a strategy that could overcome barriers to CNS drug delivery, including poor BBB penetration, extensive binding to plasma proteins, and the presence of drug efflux pumps that diminish drug levels crossing the BBB into the CNS parenchyma (Bhaskar et al. 2010; Kanwar et al. 2012b; Nowacek et al. 2009a). Various nanotechnology-based systems are being developed to overcome barriers to drug delivery for improving CNS treatments. These systems are discussed in relation to their potential use in neuroinflammatory and neurodegenerative disorders.

## 2.1 Liposomes

Liposomes are spherical vesicles composed of a lipid bilayer membrane surrounding an aqueous inner core (Fernandes et al. 2010; McMillan et al. 2011). Their high biocompatibility, low immunogenicity, and extended circulation time contribute to their use as drug-delivery systems. Hydrophilic drugs can be loaded into the aqueous core, while hydrophobic molecules can be carried in the lipid membrane for delivery to the target site. However, low loading capacity, relatively quick release profiles of hydrophobic drugs, and lesser physical stability than polymeric formulations limit their applications for specific drugs (Liu et al. 2010). Inclusion of cholesterol into the liposomal membranes can improve stability of the bilayer, coating with polyethylene glycol (PEG) chains can enhance the circulation time of liposomal formulations in vivo by reducing uptake by the reticuloendothelial system, and adding targeting moieties on the liposome surface can improve delivery to the target site (Fernandes et al. 2010; McMillan et al. 2011; Micheli et al. 2012). The mechanism by which liposomes transverse the BBB is still not understood but is proposed to be achieved by passive diffusion through the brain capillary endothelial cells, by endothelial cell endocytosis, or by fusion with the endothelial cell membrane (Fernandes et al. 2010). Improved CNS drug delivery and reduced toxicity have been observed in animal models using liposomal systems for antifungals, antiretrovirals, antiepileptics, anti-ischemia drugs, and chemotherapeutics (Wong et al. 2012). Cationic liposomes are being explored to improve delivery of nucleic acids to target cells for gene therapy (Fernandes et al. 2010).

## 2.2 Solid Lipid Nanoparticles and Nanocapsules

Other lipid-based systems that show potential for enhanced drug delivery across the BBB include solid lipid nanoparticles and lipid nanocapsules (Fernandes et al. 2010; Wong et al. 2012). These systems usually use triglycerides and phospholipids to encase the drug and are stabilized using surfactants. Solid lipid nanoparticles containing nitrendipine or 5-fluoro-2-deoxyuridine improved penetration of the drugs into the brain up to twofold (Fernandes et al. 2010; Manjunath and Venkateswarlu 2006; Wang et al. 2002). Indomethacin-loaded nanocapsules reduced tumor size and decreased morbidity and mortality in rats implanted with C6 glioma (Bernardi et al. 2009).

## 2.3 Polymeric Micelles

Polymeric micelles can be used to deliver poorly soluble drugs with high bioavailability (Gaucher et al. 2010; Kanwar et al. 2012a; Kedar et al. 2010). Polymeric micelles have an anisotropic water distribution with a hydrophilic corona and hydrophobic core. Amphiphilic block polymers containing poly (ethylene) oxidepolypropylene oxide are most commonly used in assembly of polymeric micelles for targeted drug delivery. Kabanov et al. demonstrated CNS delivery of micelles formed by polymeric block polymers surrounding a non-covalently bound drug core (Manickam et al. 2012; Sharma et al. 2008). One of the major important advantages of this system is the ability of the micellar formulations to be used in applications involving drug delivery across the BBB, since the micellar formulations are able to transport the active agents into body compartments where the free drug alone cannot penetrate (Manickam et al. 2012; Sharma et al. 2008). They also have a low critical micelle concentration and slow rate of dissociation that allows the loaded drugs to remain in the system for a longer period of time and provides greater drug delivery to the target site (Batrakova et al. 2011; Fernandes et al. 2010; McMillan et al. 2011).

## 2.4 Polymeric Nanoparticles

Polymeric nanoparticles are prepared using a variety of polymers. Intended drugs can be dissolved or encapsulated to obtain nanospheres or nanocapsules by attaching the drug to the surface of the nanoparticle (Gupta and Jain 2010; Mallipeddi and Rohan 2010). Nanocapsules are fabricated by confining the drug to a cavity surrounded by a polymer membrane, and nanospheres are matrix systems formed by a uniform dispersion of the drug (Gupta and Jain 2010). The development of biodegradable polymers and their use for various drug-delivery applications have been described recently. The most widely used nontoxic polymeric materials include polyvinyl alcohol (PVA), polyacrylic acid (PAA), polyacrylamide, PEG, and polycaprolactone (Gupta and Jain 2010). These polymers

could break down easily into biologically acceptable molecules. Polymeric nanoparticles may cross the BBB by passing through tight junctions; enhanced binding, endocytosis, and transcytosis by endothelial cells; or inhibition of endo-thelial cell efflux transporters (Fernandes et al. 2010; Kanwar et al. 2012a; Khalil et al. 2011; Wong et al. 2012).

## 2.5 Dendrimers

Dendrimers are macromolecules with a defined branched architecture. They are assembled in layers, or generations, and are composed of a series of repeated branched chains around a central core with many identical functional groups on the exterior (Boas and Heegaard 2004). Drugs or other molecules can be carried within the void spaces in the interior of the structures, while attachment of functional groups on the exterior can provide targeting to specific cells and tissues (Fernandes et al. 2010; McMillan et al. 2011). The ability to pass through the BBB is greatly influenced by the molecular architecture (Fernandes et al. 2010).

## 2.6 Fullerenes, Carbon Nanotubes, and Carbon Nanofibers

Biomedical application of fullerenes and carbon nanotubes has been limited due to their poor water solubility. Functionalization has increased their water solubility and utility for use in nanomedicine. Use of functional moieties such as dendrimers, amino acids, peptides, proteins and growth factors, liposomes, polyamines, polymers, carboxyl groups, and gadolinium has expanded their application for drug delivery, reactive oxygen species quenching, promotion of neurite outgrowth, and imaging. Carbon nanofibers have potential for design of neural prosthetics (Kanwar et al. 2012b; McMillan et al. 2011).

## 3 Nanoparticles for Imaging and Diagnostics

Nanoparticles are used for neuroimaging and are increasingly being used as tools to help identify biomarkers of disease and monitor the success of stem cell treatment (Kanwar et al. 2012b; McMillan et al. 2011; Stoll and Bendszus 2010). Liposomes, dendrimers, magnetic nanoparticles, and quantum dots are being developed for single photon emission computed tomography, positron emission tomography, magnetic resonance imaging (MRI), fluorescence microscopy, computed tomography, and ultrasound. Of importance, these nanoparticles can be targeted to specific cell types and proteins to enhance sensitivity and improve detection of early markers of disease including BBB integrity and secretion of pro-inflammatory markers (Kanwar et al. 2012b). Incorporating multiple contrast agents into a single particle can provide for multiple imaging modalities, while incorporating drug and contrast agents into one particle can enable in vivo tracking of the drug distribution (Bhaskar et al. 2010; McMillan et al. 2011). Iron particles are commonly used as contrast agents in the

form of superparamagnetic iron oxide (SPIO) or ultrasmall superparamagnetic iron oxide (USPIO) (Bhaskar et al. 2010) and provide a means to track the cells and determine CNS infiltration by MRI (Stoll and Bendszus 2010). SPIO have been used successfully in experimental models of MS, stroke, and other CNS disorders to track the migration of circulating monocytes/macrophages over the course of inflammation (Bhaskar et al. 2010; Stoll and Bendszus 2010).

## 4 Neuroimmunology and Nanomedicine

### 4.1 Innate and Adaptive Immune Responses

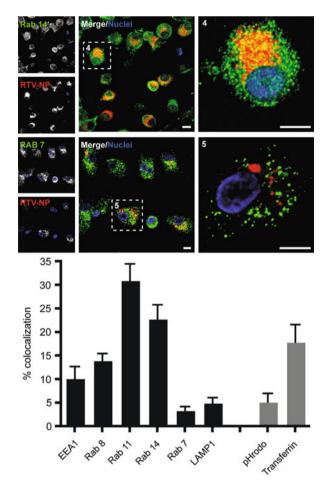
Neuroinflammation is centered on microglial activation (de Vries et al. 1997, 2012). In the presence of an activating stimulus, microglial surface receptor expression and production and secretion of immunomodulatory factors is altered (reviewed by de Vries et al. (2012) and Frank-Cannon et al. (2009)). Glial production of specific pro-inflammatory mediators such as tumor necrosis factor (TNF) and interleukins-1 (IL-1), -2, -4, -5, -8, -10, and -12 alter brain capillary endothelial cell function and promote leukocyte infiltration into the CNS (Benoit et al. 2008; de Vries et al. 1997; Persidsky et al. 2006; Ransohoff and Engelhardt 2012). Secretion of interleukins, such as IL-6 and -8, can promote both leukocyte infiltration and neuroinflammation (Benoit et al. 2008; de Vries et al. 1997; Persidsky et al. 2006). Glial production of interferon gamma (IFN- $\gamma$ ) and macrophage colony-stimulating factor (MCSF) stimulates release of eicosanoids, reactive oxygen and reactive nitrogen species, and chemokines, such as RANTES and macrophage inhibitory proteins-1 alpha and -1 beta (MIP-1 $\alpha$  and MIP-1 $\beta$ ) (de Vries et al. 1997; Persidsky et al. 2006). Infiltration of neutrophils, monocytes, and antigen-specific B and T cells into the CNS is accelerated by the expression of intercellular adhesion molecule-1 (ICAM-1), ICAM-2, and vascular cell adhesion molecule-1 (VCAM-1) on brain capillary endothelial cells (de Vries et al. 1997; Persidsky et al. 2006) and their interactions with lymphocyte functional antigen-1 (LFA-1) and very late antigen-4 (VLA-4) (de Vries et al. 1997; Kanwar et al. 2012a; Lyck and Engelhardt 2012). IL-1 beta (IL-1 $\beta$ ) is one example of the range of inflammatory factors that affects endothelial cell expression of adhesion molecules, enhances chemoattractant secretions by astrocytes, and promotes inflammatory cell migration (Brabers and Nottet 2006). Chronic activation of this system designed to fight CNS inflammation in disease or injury is believed to underlie the pathobiology of neurodegenerative diseases (de Vries et al. 1997, 2012).

## 4.2 Cell-Based Drug Delivery

The application of cells as targeted carriers for nanomedicines is a developing concept. Because of their ability to migrate to sites of disease and inflammation, immunocytes, mononuclear phagocytes (MP; monocytes, macrophages, dendritic

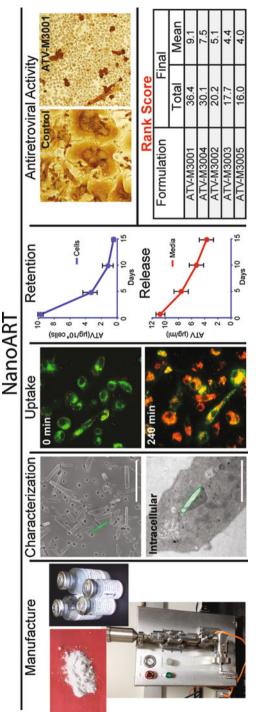
cells), lymphocytes, neutrophils, and stem cells are attractive as drug carriers (Batrakova et al. 2011; McMillan et al. 2011). They can also act as Trojan horses for delivery of drug cargo across biologic barriers such as the BBB. For cells to be effective drug carriers, however, a number of criteria must be fulfilled. First, the cells must migrate to the site of injury or disease in sufficient numbers. For CNS disorders, this means that the cells must be able to cross the BBB. Second, the cells must release their drug cargo intact at the disease site. Thus, the drug must be carried in non-degrading compartments within the cell and be released from the cell continuously at the site of action. Finally, carriage of the drug cargo should not adversely affect the cell's normal function. The potential for cell-based drug delivery has been demonstrated by our own studies. Macrophages loaded with indinavir nanoparticles and injected intravenously into severe combined immunodeficient (SCID) mice migrated to a site of HIV-encephalitis (HIVE) in the brain, reduced the level of infection, and provided enhanced drug levels at the site of infection (Dou et al. 2009). As a second example, in an experimental model of Parkinson's disease, macrophages loaded with catalase nanozymes were injected into 1-methyl 4-phenyl 1,2,3,6-tetrahydropyridine (MPTP)-treated mice and provided enhanced catalase levels in the brains of these mice (Batrakova et al. 2007; Brynskikh et al. 2010). Drug loading of neural stem cells and their use as drugdelivery vehicles to areas of CNS disease is also being explored (Martinez-Serrano and Bjorklund 1998; Martinez-Serrano et al. 1996; McMillan et al. 2011; Muller et al. 2006). The success of a nanocarrier for use in a cell-based delivery system is determined by a number of factors. Drug nanocarriers are commonly composed of a polymeric outer shell with an inner core for drug carriage. The charge state of the outer polymeric shell can directly influence the uptake of the nanoparticles by immunocytes and stem cells through cell surface receptor interactions and stability of the nanoparticles inside the cells (Batrakova et al. 2011; McMillan et al. 2011). As an example, nanozymes containing superoxide dismutase were protected inside macrophages when the nanoparticles were made using positively charged block copolymers but not negatively charged block copolymers (Batrakova et al. 2011; Zhao et al. 2011). The composition of the outer polymeric coating can also influence the toxicity of the nanoparticles to the cells (Batrakova et al. 2011; McMillan et al. 2011). We have demonstrated that nanoformulations of the antiretroviral drug ritonavir, which enter cells through clathrin-mediated endocytosis, are trafficked to and stored in recycling endosomes and can be released intact (Fig. 1) (Kadiu et al. 2011). Of particular interest these compartments are the same compartments where the human immunodeficiency virus-1 (HIV-1) replicates (Kadiu and Gendelman 2011).

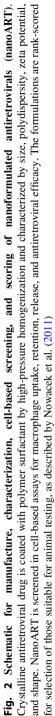
While these experiments demonstrate the potential for cell-based drug delivery, they require pre-loading of cells with nanoparticles and injection into donors. A more practical, clinically useful system would provide for administration of drug nanoparticles to the diseased patient; uptake of the nanoparticles by host MPs, immunocytes, or lymphocytes; and storage within these cells to provide (a) sustained drug release and (b) delivery of the drug nanoparticles to sites of CNS disease. We have recently demonstrated this possibility using polymeric crystalline

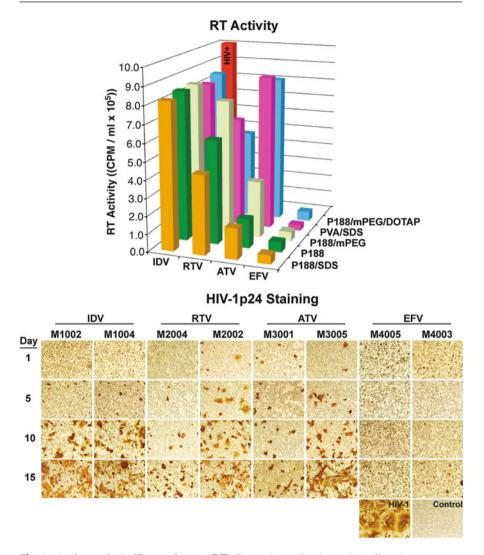


**Fig. 1** Immunohistological identification of nanoparticle subcellular localization. Confocal microscopy confirmed distribution of RTV-NP within recycling endosomal (Rab 14) rather than late endosomal (Rab 7) compartments. Note that RTV-NPs are *red* and compartments are *green*, and *yellow* signifies compartment marker-particle overlap in all panels. Pearson's co-localization coefficients indicate RTV-NPs are preferentially distributed to Rab11 and Rab14 recycling endosomes compared with early endosomes, Rab8 or Rab7 endosomes, and lysosomes. Analysis of distribution of RTV-NP within acidified (degrading) compartments, identified by pHrododextran beads, revealed minimal overlap, indicating RTV-NPs likely bypass degradation within the cell and are primarily recycled for release. High RTV-NP co-localization with transferrin also indicates that particles are most likely recycled. Measure bars equal 1 µm. Graphical data represent the mean  $\pm$  standard error of the mean for n = 3. *NP* Nanoparticle, *RTV* Ritonavir (Figure reprinted with permission from Kadiu et al. (2011))

nanoparticles of the antiretroviral drugs atazanavir (ATV) and ritonavir (RTV) (Fig. 2). Polymer-coated crystalline nanoparticles of antiretroviral drugs (nanoART) are taken up by macrophages in culture, retained inside the cells for up to 15 days, and demonstrate antiretroviral efficacy over this time period







**Fig. 3** Antiretroviral efficacy of nanoART. Comparison of antiretroviral effects in monocytederived macrophages challenged with  $\text{HIV-1}_{ADA}$  15 days after pretreatment with nanoART, as measured by reverse transcriptase (RT) activity and  $\text{HIV-1}_{P24}$  antigen staining 10 days after viral challenge. RT activities were measured by <sup>3</sup>H-TTP incorporation. Data represent mean for n = 8 determinations/treatment (Figure reprinted with permission from Nowacek et al. (2011))

(Fig. 3) (Balkundi et al. 2011; Nowacek et al. 2009b, 2010, 2011). When administered subcutaneously to mice, nanoART (250 mg/kg, ATV/RTV) provided sustained plasma ATV concentrations (>100 ng/ml) and liver and spleen levels exceeding 1,000 ng/g (Roy et al. 2012) for up to 2 weeks. Predominant sequestration of nanoART inside liver Kupffer cells was demonstrated in mice

treated with dye-labeled nanoART followed by isolation and selection of Kupffer cells by in situ collagenase digestion and CD11b + cell sorting (Gautam et al. 2013). Of significance, weekly administration of nanoART reduced HIV-1 viral load to nearly undetectable levels in a humanized mouse model of chronic HIV infection (Dash et al. 2012). Targeting the folate receptor on macrophages using a folate-modified polymer coating improved by up to twofold in vitro macrophage uptake and antiretroviral efficacy of nanoART and enhanced mouse plasma and tissue drug levels up to fivefold compared to non-targeted nanoART (Puligujja et al. 2013).

# 5 Degenerative, Inflammatory, and Infectious Diseases of the Nervous System

# 5.1 AD, PD, and ALS

#### 5.1.1 Pathobiology of Diseases

AD, PD, and ALS are progressive neurodegenerative disorders that usually develop later in life. The etiologies of these diseases are different but have in common deposition of mutated proteins and chronic neuroinflammation (Kanwar et al. 2012b; Nowacek et al. 2009a; Re et al. 2012b). AD is characterized by losses in memory and cognitive function and pathologically by neurofibrillary tangles formed by hyperphosphorylated tau protein and extracellular  $\beta$ -amyloid protein (A $\beta$ ) plaques, considered the cause of neuronal loss. PD is marked by degeneration of dopaminergic neurons in the substantia nigra and characterized pathologically by cytoplasmic inclusions called Lewy bodies composed of mutated  $\alpha$ -synuclein. ALS is marked by the selective loss of motor neurons in the brain and spinal cord leading to paralysis. The majority of ALS cases are sporadic; however, the familial form has been linked to missense mutations in superoxide dismutase 1 (SOD1), which forms intracellular deposits that inhibit chaperone and proteasome activity.

### 5.1.2 Nanoparticle-Based Therapies and Diagnostics

Approved therapies for these diseases are mainly symptomatic. AD treatments include acetylcholinesterase inhibitors to prevent breakdown of acetylcholine and the use of antioxidants and anti-inflammatory drugs (Kanwar et al. 2012a). Treatments for PD include dopamine mimetics and monoamine oxidase inhibitors to reduce dopamine degradation (Kanwar et al. 2012a). The only currently approved treatment for ALS is riluzole, a sodium channel blocker that inhibits stimulation of glutamate receptors and is used to slow the progression of the disease. None of these treatments are curative, and there is a grave need for novel therapeutic and early diagnostic approaches.

Development of nanotherapeutic-based treatments for each of these diseases is ongoing (Kanwar et al. 2012b; Nowacek et al. 2009a). However, while some are promising, none have reached clinical trials. Again, many, but not all, of the nanoparticle-based systems are designed to enhance the effectiveness and reduce the toxicity of traditional treatments. Use of nanoparticles for treating AD has

focused on three strategies: prevention of Aβ aggregation in the CNS and blood; reducing transition-metal-induced production of reactive oxygen species; and improved delivery of acetylcholinesterase inhibitors to the CNS (Nowacek et al. 2009a; Re et al. 2012a). Polyamidoamine (PAMAM) dendrimers, and nanoliposomes decorated with phosphatidic acid, cardiolipin, anti-A<sup>β</sup> antibody or a curcumin derivative can inhibit in vitro aggregation of A $\beta$ , and C<sub>60</sub> fullerenes were reported to prevent cognitive impairments induced by A $\beta$  aggregation in rats (Nowacek et al. 2009a; Re et al. 2012a). Aberrant homeostasis of transition metals in the CNS has been associated with oxidative insult to neurons and promotion of A $\beta$  aggregation. The use of metal chelators has been proposed, but is complicated by their systemic toxicity and inability to cross the BBB. To improve the therapeutic efficacy and CNS delivery of chelators and reduce their unwanted systemic toxicity, nanoparticle systems are being developed (Cui et al. 2005; Liu et al. 2009a,b). Nanoparticle systems to improve delivery of anticholinesterase inhibitors to the brain, such as polysorbate 80-coated poly(n-butylcyanoacrylate) nanoparticles to deliver tacrine and poly(DL-lactide-co-glycolic acid) (PLGA) microspheres to deliver huperzine-A, are also under development (Chu et al. 2007; Nowacek et al. 2009a; Re et al. 2012a; Wang et al. 2007; Wilson et al. 2008).

Nanoparticles for treatment of PD are being developed to treat loss of dopamine and dopaminergic neurons and reduce microglia-activated reactive oxygen species production (Nowacek et al. 2009a; Re et al. 2012a). To deliver dopamine to the CNS, chitosan nanoparticles containing dopamine adsorbed to the surface were developed and provided reduced toxicity and increased dopamine levels in the striatum (Trapani et al. 2011). Solid lipid nanoparticles containing bromocriptine, a dopamine agonist, improved performance in behavioral tests when administered to rats (Esposito et al. 2008). Lactoferrin-modified nanoparticles containing human neurotrophic factor were used for gene therapy in a rat model of PD and provided improved locomotor activity, reduced neuronal loss, and increased brain dopamine levels.

Fewer nanoparticle treatments are under development for ALS than for AD and PD; however, the approaches being used are similar (Nowacek et al. 2009a; Re et al. 2012b). Solid lipid nanoparticles containing riluzole exhibit high drug loading, a greater ability to deliver drug to the brain, and a higher efficacy compared to free drug (Bondi et al. 2010). In addition, PLGA nanoparticles containing SOD1 and carboxyfullerene SOD mimetics show protection from hydrogen-peroxide-induced damage in neuronal cultures, (Ali et al. 2008; Reddy et al. 2008) which opens the way for these systems to be tested in animal models of ALS.

A cell-based Trojan horse approach for treatment of neurodegenerative disease has been developed by Batrakova and coworkers (2007, 2011; Brynskikh et al. 2010). Catalase was packaged into nanoparticles composed of a block copolymer complex with polyethyleneimine-PEG (termed nanozymes) and loaded into mouse bone marrow macrophages in vitro. The macrophages loaded with nanozymes were adoptively transferred into isologous mice treated with MPTP as a model of acute PD. Nanozyme-loaded macrophages were able to cross the BBB and deliver active catalase to sites of damage to a greater extent that free catalase (Batrakova et al. 2007), providing proof that this concept is feasible and could be applied to other neurodegenerative disorders.

Nanotechnology also offers tools for early disease diagnosis. Ultrasensitive assay systems for detecting low concentrations of AD pathogenic proteins using gold nanoparticles have been developed by Georganopoulou et al. (2005) and Neely et al. (2009). Amyloid plaques in brains of AD transgenic mice were detected using A $\beta$ -coupled iron oxide particles and MRI (Yang et al. 2011). For early diagnosis of PD, sensitive in vitro nanoparticle-based systems are being developed for detection of  $\alpha$ -synuclein (An et al. 2010; Yu and Lyubchenko 2009). MRI has also been used to track CD4+ lymphocytes loaded with SPIO nanoparticles and injected into a rat model of ALS to detect regions of pathology in the CNS (Bataveljic et al. 2011; Machtoub et al. 2011).

# 5.2 Multiple Sclerosis

### 5.2.1 Pathobiology of Disease

MS is a progressive inflammatory autoimmune disorder of the CNS characterized by demyelination and degeneration of neuronal axons (Goldenberg 2012; Lassmann and van Horssen 2011). Destruction of oligodendrocytes occurs upon infiltration of lymphocytes into the CNS and initiation of inflammation and autoimmune reactions (Kanwar et al. 2012a). The disease can be manifest as a relapsing remitting MS, characterized by recurring attacks of acute focal neurological deficits or exacerbation of existing deficits followed gradually by partial or full recovery, or a progressive steady decline without remission (Goldenberg 2012). The disease course includes distinct exacerbations with no symptom-free remissions; instead, patients will experience progressive acceleration of disability, accumulating irreversible neurologic deficits.

The pathology of MS was originally defined by the presence of focal white matter lesions, characterized by primary demyelination with partial preservation of axons and reactive astrocytic scar formation (Bo et al. 2003; Nylander and Hafler 2012). The disease pathology changes with time. For patients in the early stages of disease experiencing clinical relapses and remissions, inflammatory demyelination leads to the formation of focal plaques predominantly located in the white matter (Rovira and Leon 2008). In latter stages of the disease, widespread demyelination in the cerebral and cerebellar cortex is seen as well as diffuse degenerative changes throughout the entire white and gray matter. Long-standing severe disease results in profound brain and spinal cord atrophy, with extreme tissue loss and dilatation of cerebral ventricles (Androdias et al. 2010; Kutzelnigg et al. 2005; Popescu et al. 2011). In a substantial number of patients, the disease eventually subsides and inflammation as well as neurodegeneration decline to levels seen in age-matched controls. During stages of active disease, both in the relapsing as well as in the progressive stages, profound oxidative tissue injury is seen. In early stages, oxidative damage is mainly driven by inflammation, resulting in an oxidative burst in microglia and macrophages. In the progressive stage, active neurodegeneration occurs against a background of mild-to-moderate inflammation, and oxidative

damage appears to be augmented by release of iron from intracellular stores. In the latter stages, the presence of iron in the brain in the absence of additional oxidative bursts from microglia does not lead to further neurodegeneration; however, patients may be affected by concomitant diseases such as stroke or AD.

### 5.2.2 Current Treatments

Traditional treatment for MS is aimed at shortening the duration and frequency of acute exacerbations and providing symptomatic relief. Most currently approved MS therapeutics are immunosuppressants with the exception of beta interferons and glatiramer acetate as disease-modifying drugs for relapsing remitting MS (Gasperini and Ruggieri 2012; Silva 2005; Tubaro et al. 2012). The mechanism by which these drugs act involves downregulation of T-cell proliferation and migration, reduction in antigen presentation, and shifting the immune response to T-helper 2 (Th2) cells (Devonshire and Verdun di Cantogno 2011). The necessity for repeated subcutaneous or intramuscular injections of these drugs can lead to suboptimal long-term adherence in some patients and can negatively affect clinical outcomes. Other treatments target immune suppression by inducing apoptosis in lymphocytes (mitoxantrone) or T and B cells (cyclophosphamide) or using corticosteroids to suppress the immune response. While these are commonly used for treatment of relapsing remitting MS, they have limited application in the treatment of other forms of MS because of undesirable side effects.

## 5.2.3 Nanoparticle-Based Experimental Systems

Nanoparticle-based treatments for MS are being developed that seek to not only modulate the immune response, but also promote neuroprotection and myelin regeneration (Amedei et al. 2012; Dharamkar 2008; Ellis-Behnke 2007; Silva 2005; Ulbrich and Lamprecht 2010). PEG-coated liposomal formulations of the traditional MS therapeutics, prednisolone and minocycline, are being developed for enhanced delivery of drug to inflammatory lesions in the CNS and to minimize serious systemic side effects (Hu et al. 2009; Linker et al. 2008; Schmidt et al. 2003a, b). PEG-coated liposomal prednisolone administered intravenously to rats with experimental autoimmune encephalomyelitis (EAE), an animal model of MS, provided a 4.5-fold higher concentration of drug in the spinal cord in comparison to rats without disease (Schmidt et al. 2003a, b). High levels of prednisolone in the CNS were observed only in EAE rats and were accompanied by low levels of free drug in the serum. In other studies Linker et al. (2008) observed that liposomal methylprednisolone and PEG-liposomal prednisolone had superior efficacy to methylprednisolone pulse therapy in EAE rats through second or third relapses. Studies by Hu et al. (2009) demonstrated that PEG-liposomal minocycline, administered every 5 days, was as effective as daily injections of non-formulated minocycline at reducing CNS manifestations in EAE rats. These results suggest that long-acting nanoformulations of glucocorticoids and minocycline can improve pharmacokinetics and drug dosing while reducing unwanted side effects in MS treatment.

Nanoparticle systems that deliver non-traditional therapeutics for MS are also being developed. Therapeutics that aid neuroprotection and neuroregeneration are of particular interest. A water-soluble fullerene derivative ABS-75 attached to an *N*-methyl-D-aspartate (NMDA) receptor antagonist was designed to contain both antioxidant (fullerene) and anti-excitotoxic (NMDA receptor antagonism) properties (Basso et al. 2008). Fullerene ABS-75 reduced the clinical progression of EAE in a mouse model immunized with myelin oligodendrocyte glycoprotein (MOG). This was accompanied by reduced axonal loss and demyelination and inhibition of oxidative injury, CD11b+ cell infiltration, and CCL2 expression in the spinal cord. Nanoparticles have also been proposed as a way to deliver myelin to demyelinated cells (Dharamkar 2008) and as scaffolds for neuroregeneration (Ellis-Behnke 2007; Silva 2005).

In addition to drug delivery, nanoparticles are being used to aid in the development of stem cell therapies for inducing local neurogenesis or myelogenesis in CNS lesions. Mesenchymal stem cells (MSCs), which can give rise to neural-, glial-, and astrocytic-like cells in vitro, were labeled with the SPIO MRI contrast agent ferumoxides (Feridex) and injected intrathecally and intravenously into MS patients (Karussis et al. 2010). MRI revealed the presence of MSCs in the occipital horns of the ventricles, indicating the possible migration of cells into the meninges, subarachnoid space, and spinal cord. In addition, an increase in CD4+ CD25+ T cells and a decrease in lymphocyte proliferation and expression of CD40+, CD83+, CD86+ and HLA-DR on myeloid dendritic cells indicated immunomodulatory effects of MSC transplantation. Further studies by Cohen et al. (2010) demonstrated that uptake of ferumoxides by neural precursor cells (NPCs) produced no adverse effects on in vitro cell survival or differentiation or in vivo functional response, suggesting that tracking of ferumoxides-labeled NPCs by MRI is a viable noninvasive method for monitoring transplantation.

A hallmark of the immune response in disorders such as MS is a deficit in the numbers of T regulatory (Treg) cells (Buckner 2010; Wing and Sakaguchi 2010). In an effort to enhance Treg numbers, PEG-coated gold nanoparticles were used to co-administer the aryl hydrocarbon receptor activator, 2-(1'H-indole-3'-carbonyl)-thiazole-4-carboxylic acid methyl ester (ITE), and a T-cell epitope from MOG (MOG<sub>35-55</sub>) to promote the generation of Tregs by dendritic cells (Yeste et al. 2012). When administered to EAE mice, the ITE-MOG<sub>35-55</sub> nanoparticles expanded the FoxP3<sup>+</sup> Treg population and suppressed the development of disease.

# 5.3 Microbial Infections

### 5.3.1 Bacterial Meningitis

## Pathobiology of Disease

Meningitis is an inflammation of the brain and spinal cord caused by chemical irritation, drug allergies, tumors, fungal infections, and infectious agents (Yogev and Tan 2011). Viral meningitis, most commonly caused by a number of enteroviruses, but also by herpes viruses and West Nile virus, occurs frequently but usually resolves without treatment. Treatment with antivirals is required for herpes virus infections (Stahl et al. 2011a, b).

Bacterial meningitis is more severe and can result in brain damage or even death if not treated. The most frequent causative agents are Hemophilus influenzae, Neisseria meningitidis and Streptococcus pneumoniae. The only clinically available vaccine is against *H. influenzae* (McIntyre et al. 2012; Schuchat et al. 1997; Yogev and Tan 2011) and has been very effective in eliminating this disease when introduced into a population (Schuchat et al. 1997; Yogev and Tan 2011). Effective antibiotics are available, although the mortality rate in the very young and the elderly remains high. However, the increasing development of antibiotic-resistant organisms is of great concern. The pathogenesis of bacterial meningitis involves the infiltration of bacteria into the subarachnoid space. Bacterial degradation products stimulate the release of pro-inflammatory cytokines, such as TNF $\alpha$  and IL-1 $\beta$ , and prostaglandins by leukocytes, microglial cells, astrocytes, and endothelial cells, leading to increased BBB permeability. Infiltration of neutrophils and leakage of plasma proteins produces further damage (de Vries et al. 1997). Antibiotics can halt bacterial replication but are not effective alone in reducing the inflammatory events that contribute to brain damage and long-term disability (Borchorst and Moller 2012; de Vries et al. 1997; van de Beek et al. 2012). Adjunctive therapy with corticosteroids to reduce inflammation is recommended, but the incidence of detrimental side effects is high (Borchorst and Moller 2012). Thus, there is a need for new therapeutics that can penetrate the BBB and deliver traditional and novel antibiotics and adjuncts to the site of infection and inflammation while reducing the incidence and severity of side effects.

Complicating effective treatment of the disease are the time-consuming methods used for diagnosis, which can delay initiation of therapy and contribute to morbidity and mortality. Current diagnostic methods use conventional microscopy and bacterial culture of blood and cerebrospinal fluid followed by confirmation and sero-type identification by polymerase chain reaction (PCR) (Bronska et al. 2006; Reddy et al. 2012; Rosenstein et al. 2001; Tunkel et al. 2004). New methods to speed diagnosis and expedite drug therapy are sorely needed.

### Nanoparticle-Based Therapies

A number of experimental nanoparticle-based systems for diagnosis and treatment of infectious meningitis have been described. A recent study by Reddy et al. (2012) described the development of an acoustic wave immunosensor for meningococcal antigen that is enhanced using gold nanoparticles. Cell surface outer membrane protein 85 (OMP85) of *N. meningitidis* was bound to gold nanoparticles and quantitated by interaction with antibody immobilized on the surface of a polyvinylidene fluoride–coated quartz crystal microbalance. Binding of OMP85 to the gold nanoparticles allowed detection of as little as 312 ng/ml OMP85 and would provide detection of antigen in blood or CSF with minimal sample preparation.

Development and efficacy of self-assembled cationic antimicrobial peptide nanoparticles has been described (Liu et al. 2009c; Wang et al. 2010) for treatment of *Staphylococcus aureus* and *Cryptococcus neoformans* meningitis in rabbits. Cholesterol-conjugated  $G_3R_6TAT$  (CG<sub>3</sub>R<sub>6</sub>TAT) self-assembled in deionized

water into cationic core/shell nanoparticles with TAT-anchored shells. The nanoparticles crossed the BBB and were equally as effective as vancomycin and amphotericin B in treating *S. aureus* and *C. neoformans* meningitis, respectively (Liu et al. 2009c; Wang et al. 2010). In addition, they did not impair liver and kidney functions or induce blood electrolyte imbalances. Of critical importance antimicrobial peptide nanoparticles show promise for treating drug-resistant bacterial, yeast, and fungal meningitis infections. Recently, a different nanoparticle-based approach to combat development of drug-resistant *S. aureus* was described by Chakraborty et al. (2012). Folic acid–tagged chitosan nanoparticles were used to effectively deliver vancomycin into a drug-resistant strain of *S. aureus* in an in vitro study and promote bacterial lysis. Using the folic acid–conjugated chitosan nanocarrier provided an effective way of delivering vancomycin across the thicker cell wall in vancomycin-resistant bacterial strains (Chakraborty et al. 2012).

Adjunctive nanoparticle-based therapies have also been described. Tsao et al. (1999) demonstrated that *Escherichia coli*–induced meningitis could be inhibited by carboxyfullerene (C60). Treatment of *E. coli*-infected mice with a water-soluble malonic acid derivative of C60 reduced mortality, decreased levels of TNF $\alpha$  and IL-1 $\beta$  in the brain, and inhibited increases in BBB permeability and neutrophil infiltration (Tsao et al. 1999).

### **5.3.2 Human Immunodeficiency Virus** Pathobiology of Disease

HIV-associated neurocognitive disorders (HAND) comprise a spectrum of disease states from clinically asymptomatic to overt cognitive and behavioral impairments to full-blown dementia (HIVE). HIVE is fueled by immune activation of CNS cells and virotoxins, including viral proteins, cytokines, and chemokines that are secreted by infected brain macrophages and microglia (Kaul 2009; Shapshak et al. 2011). Following HIV infection, the resulting early inflammation and secreted viral proteins induce a breach in the BBB integrity, allowing HIV and infected mononuclear phagocytes to invade the CNS. Virus and toxic products secreted by infected brain mononuclear phagocytes and glial cells induce neuroinflammation, neurodegeneration, and encephalopathy. The neuropathology of HIVE is characterized by neuronal loss, glial activation, presence of multinucleated giant cells, perivascular mononuclear infiltration, myelopathy, and myelin pallor. In addition to neuronal loss, aberrant sprouting, and dystrophic synaptodendritic connections, HIVE is also characterized by extensive damage to microvessels of the blood-brain barrier caused by infiltration of HIV-infected cells and inflammatory cells into the brain (Gelman et al. 2006, 2012).

The mechanism by which HIV infection leads to neuroinflammation and encephalitis is multifactorial and has been shown to involve direct viral infection of brain mononuclear phagocytes and dysfunction and injury of brain endothelial cells, astrocytes, oligodendrocytes, and neurons, induced by HIV itself and secreted viral factors (Byrd et al. 2012; Noorbakhsh et al. 2005; Pacifici et al. 2013; Silva et al. 2012). HIVE generally occurs in the late stages of HIV infection and is associated with low levels of CD4+ T cells and high plasma viral loads. It involves continued CNS inflammation and neuronal and astrocyte injury induced by the virus, secreted viral proteins, and secreted cytokines and chemokines (Pacifici et al. 2013).

### **Current Treatments**

With the advent of effective combination antiretroviral therapy (cART) (Table 1) and the longer survival of infected patients, the pathogenesis of HIVE has been transformed from a subacute, neuroinflammatory disorder with florid neurological alterations, to a chronic and protracted condition. Advances in ART drugs and treatment regimens have improved virological control and greatly increased the life expectancy of infected individuals. However, cART has not eradicated virus from protected sites, such as the brain and lymphoid tissue, nor has it eliminated the incidence of milder forms of HAND (Valcour et al. 2011). With the need for continued life-long daily doses of cART, occurrence of adverse side effects, and lack of adherence to treatment regimens, the development of resistant strains has become a cause for concern (Chen et al. 2002; Chulamokha et al. 2005; Fellay et al. 2001; Hawkins 2006; Shehu-Xhilaga et al. 2005). Thus, there is need for novel adjunctive treatments that address HIV-associated neurological deficits and improved antiretroviral therapies to reduce the incidence of side effects and simplify dosing regimens (Mahajan et al. 2012; Nowacek and Gendelman 2009).

### Nanoparticle-Based Experimental Systems

Various nanoparticle-based systems have been explored for improving antiretroviral therapy. Efavirenz (EFV)-poloxamine/poloxamer polymeric micelles were synthesized as nanocarriers via oral administration, and displayed promising physical stability and pharmacokinetics (Seremeta et al. 2013). Intranasal administration of poly(ethylene oxide)-poly(propylene oxide) polymeric micelles loaded with EFV enhanced bioavailability of the drug in the CNS fourfold and increased the relative exposure index (ratio between the area under the curve in the CNS and plasma) fivefold with respect to the same system administered intravenously (Chiappetta et al. 2013). Polybutylcyanoacrylate and methylmethacrylatesulfopropymethacrylate nanoparticles have been reported to increase delivery of ART drugs to the CNS (Mulik et al. 2012). The disadvantages of polymeric nanoparticles include residual contamination, polymer initiation, toxic monomers, and stability issues. Shibata et al. (2013) produced biodegradable combination antiretroviral PLGA nanoparticles containing EFV and lopinavir by high-pressure homogenization. These nanoparticles also efficiently inhibited in vitro HIV-1 infection and transduction. Tenofovir, a nucleotide analog reverse transcriptase inhibitor, was embedded in PLGA and chitosan nanocarriers and demonstrated sustained in vitro release, suggesting that these particles could be effective and attractive drug carriers (Belletti et al. 2012).

ARV class	ARV agent	Abbreviation	Recommended a initial therapy
Non-nucleoside reverse transcriptase nhibitors			
	Efavirenz	EFV	Yes
	Nevirapine	NVP	Yes
	Rilpivirine	RPV	Yes
	Delavirdine	DLV	No
	Etravirine	ETR	No
Nucleoside analog reverse transcriptase nhibitors			
	Emtricitabine	FTC	Yes
	Lamivudine	3TC	Yes
	Abacavir	ABC	Yes
	Zidovudine	ZDV	Yes
	Didanosine	ddI	No
	Stavudine	d4T	No
	Apricitabine	ATC	Phase III trials
Nucleotide analog reverse transcriptase nhibitor			
	Tenofovir disoproxil	TDF	Yes
	fumarate		
Protease inhibitors			
	Atazanavir	ATV	Yes
	Darunavir	DRV	Yes
	Fosamprenavir	FPV	Yes
	Lopinavir	LPV	Yes
	Saquinavir	SQV	Yes
	Ritonavir	RTV	Yes (as boost)
	Indinavir	IDV	No
	Nelfinavir	NFV	No
	Tipranavir	TPV	No
Integrase strand transfer inhibitors			
	Elvitegravir	EVG	Yes
	Raltegravir	RAL	Yes
CCR5 antagonist			
	Maraviroc	MRV	Yes
Fusion inhibitor			
	Enfuvirtide	T20	No
CYP3A4 inhibitor			
	Cobicistat	COBI	Yes (as boost)

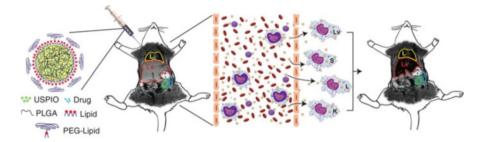
# Table 1 FDA-approved antiretroviral (ARV) drugs for HIV-1 treatment

### **Cell-Based Nanoparticle Delivery for HIV-1 Treatment**

Use of macrophages as Trojan horses to improve ART dosing by acting as drug reservoirs and providing delivery to sites of residual viral infection is being developed in our own laboratories (Balkundi et al. 2011; Nowacek et al. 2009b, 2010, 2011). We have developed polymer-coated crystalline nanoformulations of antiretrovirals (nanoART) that can be taken up by macrophages and remain inside the cells for an extended period of time (Dou et al. 2009; Nowacek et al. 2011). These formulations were manufactured by high-pressure homogenization or wet milling of crystalline drug using various surfactants including poloxamer 188 (P188), poloxamer 407 (P407), 1, 2-distearoyl-sn-glycero-3-phosphoethanolamine-N-[methoxy(polyethyleneglycol)-2000] (DSPE-mPEG), PVA, and PLGA. NanoART was efficiently and rapidly taken up by human monocyte-derived macrophages (MDM) in culture. Drug (atazanavir (ATV), ritonavir (RTV), or EFV) was retained inside the cells for up to 14 days and inhibited HIV-1 replication in macrophages in a dose-dependent manner (Balkundi et al. 2011; Nowacek et al. 2009b, 2010, 2011). NanoART-loaded macrophages could deliver drug to sites of HIV-1 infection in the brain in an animal model of HIVE. Importantly, nanoART (P188-ATV/RTV) administered subcutaneously at weekly doses (250 mg/kg each drug) provided sustained plasma and tissue ATV levels above the minimal effective concentration, suppressed viral replication in spleen and brain, and maintained CD4+ T-cell numbers in a humanized mouse model of chronic HIV-1 infection (Dash et al. 2012). These studies are an important step in the development of a cell-based delivery system for ART drugs (Balkundi et al. 2010; Nowacek et al. 2009b).

Our group has also developed nanoART targeted to macrophage receptors. Decorating nanoART with folic acid (to target macrophage folate receptors) elicited a >2-fold concentration-dependent enhancement of macrophage uptake and retention in vitro when compared to replicate non-decorated nanoART (Puligujja et al. 2013). Antiretroviral activity, measured by inhibition of HIV reverse transcriptase activity and HIV-1p24 staining, was also enhanced twofold. These activities paralleled greater nanoART retention in recycling endosomes, providing a stable cell-based drug depot. We also observed that by actively targeting the nanoART to macrophages through folate modification, plasma and tissue ATV levels could be maintained at levels above the minimum effective concentration after a single intramuscular injection (50 mg/kg) in mice (Puligujja et al. 2013).

We are also developing a novel nanoparticle system to provide a noninvasive assessment of ART tissue distribution by MRI. Small magnetite ART (SMART) particles were manufactured by encasing ATV and SPIO (magnetite) in PLGA, 1,2-distearoyl-*sn*-glycero-3-phosphocholine, and DSPE-mPEG. SMART particles were efficiently taken up by MDM in culture and retained in the cells for up to 15 days. The levels of drug and magnetite in the cells mirrored the drug/ magnetite ratio of the SMART particles (Guo et al. 2013). As shown in Fig. 4, parenteral administration of SMART particles to mice demonstrated that



**Fig. 4** Small magnetitie-containing antiretroviral (SMART) nanoparticles can be used to assess drug biodistribution. SMART particles, parentally administered to mice, are taken up by circulating monocytes/macrophages and accumulate in tissues, including liver (Lv), spleen (S), lungs (L), and kidneys (K). Magnetic resonance imaging (MRI) is used to monitor biodistribution of the magnetite-/drug-containing SMART particles (Figure reprinted with permission from Guo et al. (2013))

magnetite and drug biodistribution paralleled one another. SMART nanoparticles could facilitate the evaluation of drug biodistribution and provide rapid in vivo assessment of nanoparticles that target cell and tissue viral reservoirs. These findings can serve to improve and speed pharmacokinetic parameters in nanoparticle-based systems and to bring nanotechnology closer to human clinical application.

# 6 Nanotoxicology

Most neuroinflammatory nanomedicine studies have focused on drug delivery of nanoparticles, but the consequent adverse effects or toxicity issues of the nanocarriers cannot be ignored. Toxicity depends on various parameters of nanoparticles including morphology, size, charge, surface area, material solubility, dose, route, and duration of administration (Kanwar et al. 2012b; Liu et al. 2012). Smaller nanoparticles exhibit superior translocation kinetics; however, their smaller size could result in toxicological effects caused by the trade-off between drug potency and immunologic surveillance (Beyerle et al. 2010; Bhaskar et al. 2010; Chen and Schluesener 2008; Tian et al. 2006). The physicochemical properties of nanosized materials differ with respect to their larger counterparts, increasing the potential for toxic interactions (Nel et al. 2006, 2009). Nonbiological nanoparticulate carriers larger than 200 nm can cause cell membrane ruffling, cytoskeletal rearrangement and stimulate endocytic machinery, promoting uptake by phagocytic cells (Harding and Song 1994; Paulo et al. 2011; Pichai and Ferguson 2012). Furthermore, some studies show that nanoparticles can be pro-oxidant and pro-inflammatory (Stoeger et al. 2006, 2009). Certain types of nanoparticles may stimulate the release of cytokines and chemokines such as IL-6, IL-1 $\beta$ , TNF $\alpha$ , reactive oxygen species, and C-reactive

protein with subsequent activation of mitogen-activated protein kinase (MAPK), redox-activated transcription factors, the transcription factor nuclear factor kappa B (NFkB), and activating protein-1 (AP-1). Long-term exposure to nanoparticles may thus result in chronic inflammation that could accelerate atherosclerosis (Fernandes et al. 2010). Data are limited, however, on the potential toxicities of long-term nanoparticle exposure and possible neurological consequences of these drug-delivery systems (Bhaskar et al. 2010; Kanwar et al. 2012b).

Although nanomedicines for treatment of neuroinflammation have shown promise, further studies on the safety of these systems are needed. To reduce the potential for adverse effects, a couple of different strategies are being used in the development of nanoparticle drug carrier systems. Novel biodegradable or biocompatible polymers are being used in the preparation of nanoparticles to reduce their toxicity and allow them to escape the reticuloendothelial system, as well as to aid drug release. Biocompatible polymers such as polylactic acid, polybutylcyanoacrylate, and polycaprolactone are being used in the preparation of nanomedicines to decrease toxicity (Kanwar et al. 2012b; Nel et al. 2006). In addition, surface functional modification of nanomedicines to actively target nanoparticles to sites of disease is being increasingly used to decrease nonspecific interactions and toxicities (Koo et al. 2005; McMillan et al. 2011).

# 7 Conclusion

Treatments of neurodegenerative and neuroinflammatory disorders require drugs acting on the CNS. Thus, development of novel nanoparticle systems for drug delivery across the BBB is a promising and urgent industry aimed at improving disease outcomes (Bhaskar et al. 2010; Fernandes et al. 2010; Kanwar et al. 2012a, b; Re et al. 2012b). Nanoparticles may be employed as drug carrier systems that assist in therapeutic targeting to the CNS. Multifunctional nanoparticles allow delivery of pharmaceutical agents into the brain and provide for neuroimaging of cell dynamics and tracking of nanoparticle delivery. Such nanocarriers and/or their combination with other drugs will drive the search for targeting specific areas in the brain and thus enhance therapies. Nanomedicine has yet to make its mark in clinical studies; however, nanoparticle-based drugs are gaining considerable interest in the pharmaceutical industry. The translation of nanoparticles for clinical use in therapeutic and diagnostic applications is promising amid recent developments. However, even though nanoparticle-based diagnostic and therapeutic agents for neuroinflammatory diseases have shown great potential, there continues to be concerns about the safety of these particles. Thus, it is also necessary to gain a better understanding of the potential neurotoxicity associated with nanoparticle-based therapies.

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# Neurodegenerative Aspects of Multiple System Atrophy

# Rowan Radford, Mathew Wong, and Dean L. Pountney

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## Abstract

Multiple system atrophy (MSA) closely resembles Parkinson's disease clinically but with a range of autonomic signs in addition to motor symptoms, resulting in its designation as a Parkinson's-plus disease/atypical parkinsonism. However,

R. Radford • M. Wong

Griffith Health Institute, Griffith University, Gold Coast, QLD, Australia e-mail: rowan.radford@griffithuni.edu.au; mathew.wong@griffithuni.edu.au

D.L. Pountney (🖂) School of Medical Science and Griffith Health Institute, Griffith University, Gold Coast, QLD, Australia

e-mail: d.pountney@griffith.edu.au

unlike Parkinson's disease that displays primarily neuronal pathology, the neurodegenerative aspects of MSA reflect widespread glial cell dysfunction and the occurrence of intracytoplasmic inclusion bodies predominantly in oligodendrocytes. Alpha-synuclein, a key protein component of the glial cytoplasmic inclusions, is thought to mediate cytotoxicity via a range of abnormal modified and aggregated soluble molecular species that may even propagate from cell to cell as cargo in exosomes. Moreover, cellular model studies suggest that the sequestration of abnormal alpha-synuclein within microscopically visible intracellular inclusion bodies occurs as a defensive response that utilizes the aggresome machinery. Although aggregation of alpha-synuclein plays a central role in the disease pathology, it is becoming clear that there is a complex interplay of factors, including heat shock proteins, inflammation, calcium homeostasis, the ubiquitin proteasome system, autophagy, and oxidative stress. Recent studies have highlighted the involvement of metallothioneins, small ubiquitin-like modifier (SUMO), and  $p25\alpha$ , thus providing several new potential targets for therapeutic intervention. Inflammatory pathways may also be exploited in new treatment approaches, although it remains to be determined whether the astrocyte and microglial activation present is secondary to alpha-synuclein aggregation or primary to the disease pathogenesis, and may even have some beneficial consequences.

## Keywords

Aggresome • Alpha-synuclein • Autophagy • Calcium • Chaperones • Metallothionein • Multiple system atrophy • Neuroinflammation • SUMO

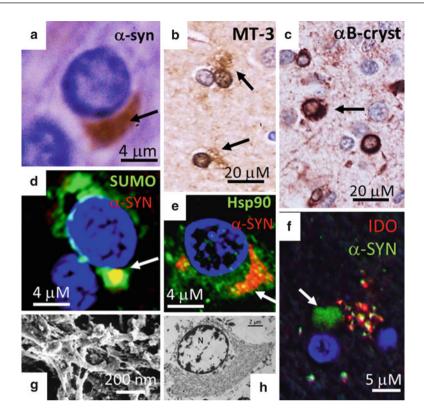
# 1 Introduction

Multiple system atrophy (MSA) is a sporadic neurodegenerative disorder clinically characterized by autonomic failure with motor syndrome presenting as atypical parkinsonism and/or ataxia. The prevalence of MSA is estimated to be 4.6 cases per 100,000 people (Marpillat 2013; Schrag et al. 1999). The prognosis of MSA is poor with death occurring within 8 years after onset (Wenning and Stefanova 2009). MSA is a heterogeneous group of neurological diseases, often misdiagnosed clinically as Parkinson's disease, which have as a common neuropathological hallmark the occurrence of filamentous, intracytoplasmic inclusion bodies in oligodendrocytes throughout the brain (Graham and Oppenheimer 1969; Papp et al. 1989; Gilman et al. 2008). Previously known as three separate clinical conditions, namely, olivopontocerebellar atrophy, striatonigral degeneration, and Shy-Drager syndrome, the signs and symptoms of MSA include autonomic malfunction, highlighted by postural hypotension, urinary complications (e.g., inability to drain bladder, urge incontinence), and persistent constipation and motor complications, such as parkinsonism, cerebellar ataxia (lack of muscle coordination), and pyramidal symptoms (Wenning et al. 2004, 2008; Wenning and Jellinger 2005; Stefanova et al. 2009). There are two subtypes of MSA as determined by the motor complication: MSA-P refers to parkinsonism, and MSA-C refers to cerebellar ataxia. In the western world, MSA-P has been shown to be the more common of the two at around 80 % prevalence (Wenning et al. 2008), although autonomic failure is seen in both subtypes (Wenning and Stefanova 2009). The macroscopic changes in brain anatomy associated with MSA have been reviewed recently (Ahmed et al. 2012). To summarize, the clinical signs, including ataxia and bradykinesia, correspond to the loss of neurons in multiple brain regions, including the striatum, substantia nigra pars compacta, locus coeruleus, cerebellum, pontine nuclei, inferior olives, and intermediolateral columns (Wenning et al. 1997; Jellinger 2003; Ozawa et al. 2004; Stefanova et al. 2009), and the severity of disease symptoms correlates well with the frequency of inclusion body-bearing cells (Papp and Lantos 1994; Jellinger and Lantos 2010). Another important aspect of the pathology of MSA is widespread gliosis associated with neuronal loss (Ishizawa et al. 2004: Stefanova et al. 2007; Song et al. 2009; Fellner et al. 2011) and the accompanying upregulation of neurotrophic factors, such as BDNF, GDNF, and IGF-I (Kawamoto et al. 1999, 2000; Pellecchia et al. 2010). The focus of this chapter is the molecular pathology of MSA, with the aim of highlighting key molecular components and their potential roles in the progression of the disease.

# 1.1 Glial Cytoplasmic Inclusions and the Aggresome Model

Many neurodegenerative diseases are characterized by microscopically visible intranuclear or intracytoplasmic protein inclusion bodies (IB), within various diseasespecific neural cell populations. In MSA, the pathological inclusion bodies occur predominantly in the oligodendrocyte cells leading to MSA being referred to as a glial  $\alpha$ -synucleinopathy, with the characteristic glial cytoplasmic inclusions (GCIs) composed largely of aggregated filamentous  $\alpha$ -synuclein (Gai et al. 1998; Wenning et al. 2008), although a subpopulation of GCIs were also immunopositive for the tau protein associated with Alzheimer's disease (Cairns et al. 1997). Figure 1a represents a typical  $\alpha$ -synuclein-positive GCI in an oligodendrocyte in MSA tissue (visual cortex). Although the inclusion bodies are seen mostly in oligodendrocytes, cytoplasmic and intranuclear neuronal inclusions and astrocyte inclusions are also seen on occasion (Nishie et al. 2004). The GCIs are often sickle shaped and found in the cytoplasm adjacent to the nucleus (Arima et al. 1992), with the inclusion body adjacent to and/or wrapped around the nucleus, and recent studies have shown that GCIs occur in mature oligodendrocytes and not in oligodendrocyte precursor cells (Ahmed et al. 2013).

In general, pathological inclusion bodies, and by extension GCIs, are believed to be formed actively in a defensive response to soluble cytotoxic protein aggregates, resembling aggresomes (Kopito 2000; Olzmann et al. 2008; Chin et al. 2010). An aggresome is formed when misfolded proteins are either not refolded with the aid of chaperones or not degraded by the normal proteolytic mechanisms (such as lysosomal autophagy and the ubiquitin-proteosome system (UPS)) (Wong and Cuervo 2010; Johnston et al. 1998). Protein misfolding can be caused by genetic or environmental factors, such as oxidative stress and metal binding (reviewed in Jomova et al. 2010). Perinuclear aggresomes are rich in aggregation-prone protein, such as  $\alpha$ -synuclein and



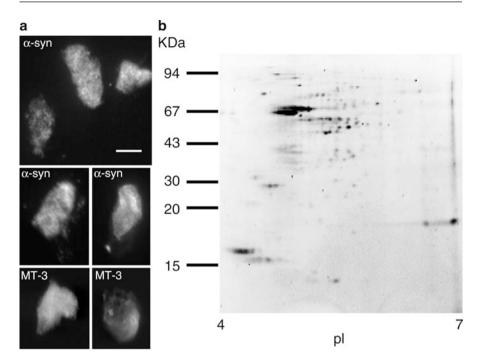
**Fig. 1** *Glial cytoplasmic inclusion bodies (GCIs) and gliosis are the primary pathological hallmarks of MSA*. Although composed largely of filamentous  $\alpha$ -synuclein aggregates, GCIs are immunopositive for ubiquitin and a diverse range of other components linked to cellular systems, such as autophagy, the proteasome pathway, inflammation, and the cytoskeleton. Illustrative micrographs of MSA brain tissue sections are presented showing (a)  $\alpha$ -synuclein, (b) metallothionein-III, and (c)  $\alpha$ B-crystallin immunostaining (visual cortex). Confocal immunofluorescence (d–f) shows the relationships between  $\alpha$ -synuclein and (d) SUMO-1, (e) Hsp90, and (f) indoleamine-2,3-dioxygenase (IDO). Panels (g–h) represent the ultrastructural features of GCIs. (g) Example scanning electron micrograph of isolated GCI surface showing  $\sim$ 10 nm single filaments and  $\sim$ 30–50 nm bundles of filaments. (h) Transmission electron micrograph of typical perinuclear GCI, showing the partial alignment of filaments towards the poles of the crescent-shaped inclusion body

ubiquitin, and cytoskeletal proteins, such as vimentin, that form a "cage" around the developing aggresome. Protein aggregates are actively transported to the microtubuleorganizing center (MTOC) along microtubules by the dynein motor proteins, along with other components to form a microscopically visible aggregate, or aggresome (Iwata et al. 2005; Olzmann et al. 2007). p97/VCP protein escorts ubiquitinated proteins to the aggresome via interactions with other ubiquitin-binding proteins, such as rad23 or HDAC6, respectively (Ju et al. 2008). This process can also be mediated by Hsp90-dependent trafficking and is inhibited by the overexpression of dynamitin that dissociates dynein from its cargo and by inhibition of tubulin re-polymerization by nocodazole (Taylor et al. 2003; Thomas et al. 2006). Recent studies have found that the parkin ubiquitin-protein isopeptide ligase (or E3) and lysine residue 63 (K63)-linked polyubiquitination are vital for dynein transport and therefore aggresome formation (Olzmann et al. 2007; Chin and Olzmann 2010). Under proteasomal impairment, parkin cooperates with the E2 enzyme Ubc13/Uev1a and PACRG to mediate Lys63-linked polyubiquitination of misfolded proteins (Taylor et al. 2012). Lys63-linked polyubiquitin chains promote binding to HDAC6 adaptor protein and thereby link the misfolded proteins to the dynein motor complex for retrograde transport towards the MTOC to form the aggresome. Binding is strengthened by editing of the K63-linked ubiquitin chains by ataxin-3 that creates free ubiquitin C-termini with increased affinity for HDAC6 (Ouyang et al. 2012). Recruitment of polyubiquitinated proteins to the aggresome is promoted by the TRIM50 protein, Lys63-linked polyubiquitination may also promote binding to the p62 protein and thereby facilitate the recruitment of autophagic membrane to the aggresome via interaction with LC3 to form an autophagosome (Fusco et al. 2012). Fusion of the autophagosome with the lysosome may allow degradation of aggregated proteins by lysosomal proteases, a process that may be overwhelmed in protein misfolding disease.

GCIs, in common with other IBs, often contain chaperone proteins, including Hsp90, and ubiquitin, which has been shown to be intimately linked to aggresome biogenesis (Olzmann et al. 2007). The GCIs are comprised largely of 9-10 nm filaments of the  $\alpha$ -synuclein protein, which is also found in the neuronal inclusion bodies (Lewy bodies) associated with Parkinson's disease and Lewy body dementia (Gai et al. 1998; Spillantini and Goedert 2000; Pountney et al. 2005a), leading to the grouping together of these diseases as  $\alpha$ -synucleinopathies. Some studies implicate the dysregulation of metals, such as copper, iron, calcium, and zinc, which are able to interact with  $\alpha$ -synuclein, inducing aggregation, oligomerization, and conformational changes in the protein (Breydo et al. 2012). Other IB components that have been identified besides  $\alpha$ -synuclein aggregates are microtubule proteins,  $\alpha$ Bcrystallin, neurofilament proteins, lipids, organelles (lysosomes and mitochondria), microtubule-associated protein 1B, and parts of the UPS (Gai et al. 1999; Ruipérez et al. 2010). Figure 1b-e illustrates the typical pattern of immunoreactivity of several GCI components in MSA tissue. Isolated GCIs can be resolved to reveal many distinct protein species (Fig. 2). An extensive list of GCI components that have been identified has been reported in the recent review by Jellinger and Lantos (2010). In the remainder of this chapter, several key factors linked to cellular systems will be highlighted that may be important to disease pathogenesis.

# 2 α-Synuclein

The main component of GCIs in MSA is aggregated filamentous  $\alpha$ -synuclein, leading to the classification of the disease as an  $\alpha$ -synucleinopathy (Gai et al. 1998; Spillantini and Goedert 2000). Indeed, GCIs containing  $\alpha$ -synuclein are required to be present upon autopsy in order for the diagnosis of MSA to be confirmed (Gilman et al. 2008). Although the origin of  $\alpha$ -synuclein in oligodendrocytes in MSA is unclear, the



**Fig. 2** Isolated GCIs show differential distributions of inclusion body components and resolve by electrophoresis to reveal abundant distinct protein species. (a) Immunofluorescence of GCIs immunoisolated from fresh, frozen MSA brain tissue using magnetic beads (Pountney et al. 2011).  $\alpha$ -Synuclein and metallothionein-III show different patterns of immunoreactivity. Scale bar, 2 µm. (b) Two-dimensional gel electrophoresis of purified GCIs (SYPRO Ruby stain, inverted) illustrates the huge diversity of distinct protein species present in the inclusion bodies (for conditions, see Pountney et al. 2005b)

accumulation of  $\alpha$ -synuclein-positive GCIs in white matter in MSA brain tissue has been shown to correlate well with the severity of tissue injury (Ishizawa et al. 2008). The  $\alpha$ -synuclein protein is encoded by the SNCA gene and comes from a family of synuclein proteins. This group encompasses three other proteins:  $\beta$ -syn,  $\gamma$ -syn, and synoretin, all of which are expressed in neuronal cells (Lavedan 1998). A wide range of in vitro and in vivo data strongly suggests a central role of  $\alpha$ -synuclein molecular interactions in the pathogenesis of  $\alpha$ -synucleinopathies (Polymeropoulos et al. 1997; Lesage and Brice 2009). A53T, G51D, E46K, and A30P mutations linked to familial PD promote the formation of soluble  $\alpha$ -synuclein protofibrils, although only the A53T and E46K mutants accelerate filament formation (Conway et al. 2000; Fredenburg et al. 2007; Kiely et al. 2013). Unlike other  $\alpha$ -synucleinopathies, such as Parkinson's disease and dementia with Lewy bodies, there have been no familial forms of MSA reported that are linked to  $\alpha$ -synuclein gene mutations, although GCI pathology has recently been found in Parkinson's disease with the G51D mutation (Kiely et al. 2013). Indeed, no clear genetic links to MSA have so far been elucidated (as reviewed by Stemberger et al. 2011; Ahmed et al. 2012), with the exception of recent studies associating MSA cases with the H1 haplotype of MAPT, commonly associated with tauopathies, such as progressive supranuclear palsy (Vilariño-Güell et al. 2011) and COQ2 mutations linked to coenzyme Q<sub>10</sub> deficiency (Tsuji et al. 2013).

 $\alpha$ -Synuclein is a protein composed of 140 amino acids, with three distinct domains: a hydrophobic central domain, a negatively charged carboxyl terminus capable of calcium binding, and an amino terminal lipid-binding domain (Ruipérez et al. 2010; Pountney et al. 2005a; Nielsen et al. 2001). Currently, the role of  $\alpha$ -syn is unknown. It is expressed in copious amounts in neurons of the central nervous system and is concentrated at presynaptic terminals near synaptic vesicles (Cookson 2009; Leong et al. 2009). In animal models, overexpression of  $\alpha$ -synuclein has been attributed to the formation of aggregates and cytotoxicity (Fernagut and Chesselet 2004; Feany and Bender 2000), and cells overexpressing  $\alpha$ -synuclein (wild-type or mutants) form  $\alpha$ -synuclein-positive cytoplasmic aggregates (Bodner et al. 2006). Indeed, oligodendrocyte expression of  $\alpha$ -synuclein in transgenic mice in conjunction with oxidative stress could replicate MSA pathology (Stefanova et al. 2005). Although, the normal cellular function of  $\alpha$ -synuclein is still not fully understood, it has been implicated in neurotransmitter vesicle binding and recycling (Nemani et al. 2010). While the free  $\alpha$ -synuclein monomer is a natively unfolded protein in dilute solution (Uversky 2009), interaction with lipid vesicles results in folding, yielding a more ordered structure (Uversky and Eliezer 2009; Perlmutter et al. 2009; Drescher et al. 2010). Many different exogenous factors, including oxidation, metal ions, and lipids, have been shown to induce the formation of potentially cytotoxic soluble  $\alpha$ -synuclein oligomers (Breydo et al. 2012).

There is no endogenous expression of  $\alpha$ -synuclein in oligodendrocytes in either MSA or normal human brain (Miller et al. 2005). Moreover, the expression of  $\alpha$ -synuclein in oligodendrocytes in rat has been found to be developmentally regulated, with no expression in mature cells (Richter-Landsberg et al. 2000). Thus, the origin of  $\alpha$ -synuclein in GCIs is likely ectopic and may result from uptake of extracellular protein. Indeed, extracellular  $\alpha$ -synuclein has recently been identified as a potential therapeutic target in  $\alpha$ -synucleinopathies (Vekrellis and Stefanis 2012). Neuronal secretion of  $\alpha$ -synuclein can occur via exosomes (Marques and Outeiro 2012), and oligodendrocyte uptake of extracellular  $\alpha$ -synuclein is mediated by endocytosis (Konno et al. 2012). There has also been much recent interest in blood and CSF  $\alpha$ -synuclein and the possibility that circulating forms of the protein could be analyzed to provide diagnostic information for  $\alpha$ -synucleinopathy diseases, including MSA (Eller and Williams 2011).

# 2.1 Alpha-Synuclein Aggregation and p25 $\alpha$

The native structure of  $\alpha$ -synuclein is a soluble unfolded or disordered conformation that can form insoluble fibrils (Vilar et al. 2008). However, the N-terminus can bind to lipid membranes where it becomes  $\alpha$ -helical in its secondary structure (Perlmutter et al. 2009; Sulzer et al. 2010). This has led to the implication that the protein may function in controlling synaptic transmission, through binding to synaptic vesicles and controlling neurotransmitter release and vesicle salvage (Nemani et al. 2010). However, studies with knockout  $\alpha$ -synuclein mice showed no detrimental signs, suggesting there are alternate methods in the control of synaptic transmission available and that  $\alpha$ -synuclein is not critical to neural development (Fernagut and Tison 2012). In vitro studies have shown that  $\alpha$ -synuclein readily forms aggregates, including both filamentous/fibrillar and oligomeric forms. Soluble oligomers of  $\alpha$ -synuclein have been shown to have a number of toxic effects, such as inhibiting the UPS, forming pores in lipid membranes, and disturbing intracellular transport (Pountney et al. 2005a; Schmidt et al. 2011). The protein C-terminus has also been shown to have a low affinity for and is required for interaction with calcium ions (Lowe et al. 2004; Nath et al. 2011), and calcium (II) binding to  $\alpha$ -synuclein has recently been shown to induce the formation of annular oligomers (Lowe et al. 2004; Nath et al. 2011).

In GCIs, ultrastructural studies have found that  $\alpha$ -synuclein is predominantly fibrillar, comprising single ~10 nm amyloid-like filaments as well as ~30 nm bundles of filaments (Gai et al. 2003), although there is also some interstitial, amorphous  $\alpha$ -synuclein aggregate. Figure 1g shows a scanning electron micrograph of the surface features of a typical GCI isolated from MSA brain tissue homogenate (Pountney et al. 2003). Furthermore, transmission electron microscopy has revealed that  $\alpha$ -synuclein filaments often align within GCIs, especially near the poles of the most common type of crescent-shaped inclusion bodies (Fig. 1h), and has also provided evidence of entrapped vesicular structures, such as mitochondria (Gai et al. 2003; Arima et al. 1992). By treating isolated GCIs with mild detergents, it was possible to release stable annular  $\alpha$ -synuclein oligomeric species (Pountney et al. 2004) that closely resemble annular  $\alpha$ -synuclein species formed in vitro (Lowe et al. 2004; Nath et al. 2011) and bacterial pore-forming toxins, such as hemolysin.

Various factors, including calcium and other metal ions (Lowe et al. 2004; Nath et al. 2011; Uversky 2009; Uversky and Eliezer 2009; Follett et al. 2012), lipids (Riedel et al. 2011), and oxidative stress (Riedel et al. 2010; Goodwin et al. 2012), have been implicated in promoting  $\alpha$ -synuclein aggregation, and recent studies have also implicated the p25 $\alpha$  protein (also known as tubulin polymerization-promoting protein, TPPP) that is differentially expressed in MSA brain tissue and itself a GCI component, in promoting the formation of cytotoxic  $\alpha$ -synuclein aggregates (Lindersson et al. 2005; Song et al. 2007; Hasegawa et al. 2010; Kragh et al. 2013).

### 2.2 Posttranslational Modifications of Alpha-Synuclein

Several modified forms of  $\alpha$ -synuclein have been reported in MSA and  $\alpha$ -synucleinopathy pathology, including Ser129-phosphorylation (Fujiwara et al. 2002; Braithwaite et al. 2012; Anderson et al. 2006; Kragh et al. 2009), glycosylation (Shimura et al. 2001), nitrotyrosine (Beyer and Ariza 2013), and various truncated and cross-linked forms of the protein (Anderson et al. 2006; Pountney et al. 2004). Recently, SUMOylation of  $\alpha$ -synuclein has been shown to inhibit filament formation and reduce neuronal toxicity in animal models (Krumova et al. 2011).

The relative importance of these different species in relation to the disease progression cannot currently be estimated; however, many recent studies have focussed on the abundant phosphorylated (phospho-Ser129)  $\alpha$ -synuclein species that can be produced by leucine-rich repeat kinase 2 (LRRK2; Qing et al. 2009), also a GCI component (Huang et al. 2008). Earlier studies have linked the glycosylated 22KDa  $\alpha$ -synuclein form to targeting of the protein to the ubiquitin proteasome pathway (Shimura et al. 2001). The nitrotyrosine modification in  $\alpha$ -synuclein has been proposed as a specific molecular marker indicating oxidative stress (Schildknecht et al. 2013). Oxidative dityrosine cross-linking has also been implicated in stabilizing  $\alpha$ -synuclein oligomers (Pountney et al. 2004). The various low-molecular weight truncated forms of  $\alpha$ -synuclein, especially the C-terminal truncations that are more prone to aggregation, have received considerable recent attention in proteomics analyses (Anderson et al. 2006), whereas the nature of the wide array of high-molecular weight, stable and cross-linked oligomeric  $\alpha$ -synuclein species is still relatively unclear (Deleersnijder et al. 2013).

# 3 Heat Shock Proteins

## 3.1 αB-Crystallin and Small Heat Shock Proteins

Heat shock proteins (Hsp), including members of the small heat shock family, have been linked to the formation of aggresomes, cytoplasmic accumulations of aggregated proteins at the microtubule-organizing center (MTOC), in response to unfolded protein stress (Berke and Paulson 2003; Bolhuis and Richter-Landsberg 2010). The chaperone activity of  $\alpha$ B-crystallin and several other small chaperone proteins, such as HSP27, is influenced by a wide range of posttranslational modifications, including phosphorylation (Aquilina et al. 2004; Kostenko and Moens 2009).  $\alpha$ B-Crystallin is upregulated in neurodegenerative diseases (Muchowski and Wacker 2005) and  $\alpha$ Bcrystallin has been shown to be present in GCIs (Gai et al. 1999). The interaction between  $\alpha$ B-crystallin and  $\alpha$ -synuclein can result in inhibition of chaperone activity (Rekas et al. 2004), and recent studies have shown that  $\alpha$ B-crystallin can influence the  $\alpha$ -synuclein fibrillation pathway (Waudby et al. 2010) and have implicated hyperphosphorylated  $\alpha$ B-crystallin forms in MSA (Pountney et al. 2005b).

# 3.2 Heat Shock Protein 90

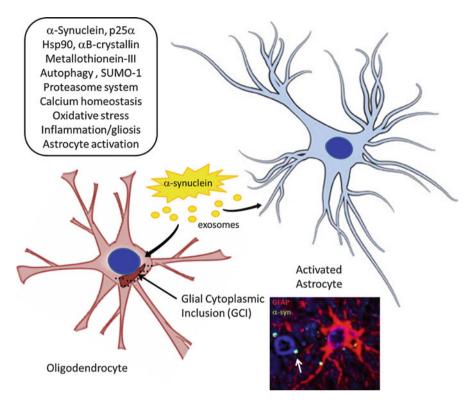
Hsp90 is a multifunctional protective chaperone protein, often working in concert with Hsp70 that is involved in the folding, refolding, and disaggregation of a diverse range of proteins. Other important functions of Hsp90 include the maintenance of tertiary structure and ATPase activity of proteasomes, activation of protein kinases, and activation of heat shock factor-1 (HSF-1). This functional diversity and specificity of Hsp90 is made possible by its conformational flexibility and dynamic association with various co-chaperone complexes. The association of Hsp70 and Hsc70 with GCIs has been reported by Kawamoto and co-workers (2007), and the association of Hsp90 with GCIs has been reported by Trojanowksi and co-workers (Uryu et al. 2006), showing granular cytoplasmic Hsp90 staining of GCI-bearing cells. Recent studies indicate that pharmacological inhibition of Hsp90 can induce Hsp70 and promote clearance and detoxification of  $\alpha$ -synuclein aggregates (Putcha et al. 2010). Hsp90 has also been linked to one form of chaperone-mediated autophagy (Kaushik et al. 2011), and the endoplasmic reticulum analogue of Hsp90 participates in the sorting of cargo proteins in autophagic clearance (Suntharalingam et al. 2012). Other recent studies have shown that Hsp90 associates with lysosomes and SUMO-1 in GCI-bearing cells in MSA (Wong et al. 2013).

# 4 Inflammation and Inflammatory Markers

Severe neuronal loss in MSA is seen in the substantia nigra, cerebellum, putamen, and locus coeruleus (Wenning et al. 1997), with neuronal loss in the brainstem and hypothalamus correlated with the autonomic dysfunction typical of MSA cases (Ozawa 2006). Considerable neuronal loss is also apparent in cortical areas, especially the motor cortex (Spargo et al. 1996; Su et al. 2001). Accompanying the neuronal loss, there is also commonly widespread demyelination observed in the white matter tracts (Papp et al. 1989). Astrogliosis and activated microglia are also found in those brain regions affected by neuronal loss and where there is a high GCI burden (Ozawa et al. 2004; Ishizawa et al. 2008), suggesting that inflammatory pathways are of major importance in MSA pathogenesis.

# 4.1 Microglia and Astroglia

The appearance of highly stellate, swollen, strongly GFAP-positive astrocytes is one of the most striking pathological features of MSA brain tissue, with this type of activated astrocyte morphology most often occurring in affected brain regions in cells neighboring GCI-positive oligodendrocytes. Figure 3 illustrates the juxtaposition of factors influencing oligodendrocyte GCI formation and astrocyte activation in MSA, with inset the typical morphology of an activated, GFAP-stained astrocyte and nearby GCI-positive oligodendrocyte observed in MSA brain tissue. The rounded morphology of activated microglia is also seen to accompany the more obvious astrocytic changes. Microglia have also been found to be more abundant in degraded regions of white matter in MSA (Ishizawa et al. 2008). The morphological indications of glial involvement in MSA is also reflected in the expression of markers for activated glia, such as increased astrocytic expression of metallothionein-III (Pountney et al. 2011) and indoleamine-2,3-dioxygenase (IDO; Radford et al. (unpublished data); Fig. 1f). Mouse models of MSA have also consistently implicated gliosis, especially microgliosis (Stefanova et al. 2005, 2007), and the interaction between  $\alpha$ -synuclein and glial cell activation has recently been linked to the TLR4 receptor (Fellner et al. 2013).



**Fig. 3** Diverse factors implicated in MSA pathogenesis result in oligodendroglial cytoplasmic inclusion body (GCI) formation and astrocyte and microglial activation. GCI formation and oligodendrocyte dysfunction together with the activation of neighboring astrocytes and microglia are the primary features characterizing MSA, leading to the secondary neuronal degeneration.  $\alpha$ -Synuclein secreted via the exosome pathway is taken up by oligodendrocytes and may act on astrocytes; however, the relationship between GCI formation and astrocyte activation remains unclear. Inset shows immunofluorescence of MSA tissue (visual cortex) stained for the astrocyte marker, GFAP, and  $\alpha$ -synuclein, illustrating a GCI-bearing oligodendrocyte (*arrow*) and the characteristic morphology of a nearby activated astrocyte

# 4.2 Metallothionein

The metallothioneins (MTs) are small (6–7 KDa) proteins, the expression of which is induced by a wide range of factors, including heavy metals, glucocorticoids, cytokines, and oxidative stress, that current evidence suggest may have a number of specific functions in the brain (Chung et al. 2008; West et al. 2008; Vasák 2005). MTs bind zinc and copper, as well as toxic heavy metals, and have also been shown to efficiently scavenge reactive oxygen species, such as nitric oxide, thereby playing a protective role against oxidative cell damage. Mammalian MT isoforms are grouped according to their charge. MT-I/II isoforms are expressed in all mammalian tissues, whereas MT-III is expressed primarily in brain and MT-IV in

squamous endothelial tissues. Recently, a significant increase was found in the number of MT-III-positive cells, especially astrocytes, in MSA brain, compared to normal brain tissue (Pountney et al. 2011). MT-III immunoreactivity was also detected in GCIs, whereas the MT-I/II isoforms were not detected, indicating a specific association between MT-III and MSA. Recent evidence shows that MT-I/II is released from astrocytes during brain injury and interacts with a neuronal cell-surface receptor to promote neuronal survival, although much less is known about the function of other isoforms. MT-III (or neuronal growth inhibitory factor, GIF) can inhibit neuritic sprouting of cultured neurons and has been reported to be downregulated in Alzheimer's disease neurons (Uchida et al. 1991; Su et al. 2001) yet is unchanged in Alzheimer's mouse models (Carrasco et al. 2006). MT-III may be linked to the cellular response to mild CNS insult, such as neurotrauma, kainic acid-induced seizures, and autoimmune responses, being coordinated with the action of various neurotrophic factors. The origin of metallothionein-III in MSA may relate to the known association of these proteins with neuroinflammation and oxidative stress (Manso et al. 2011) or may indicate a disturbance of metal ion homeostasis. MT-III has also been linked to lysosome function in astrocytes (Lee and Koh 2010). Interestingly, recent studies have shown that astrocyte expression of MT-III can promote actin polymerization (Lee et al. 2011), and so it may be that increased expression of this protein occurs as a consequence of the cytoskeletal changes during astrocyte activation.

# 5 Calcium Homeostasis

It has been speculated that calcium may play a crucial role in the pathogenesis of neurodegenerative diseases, such as  $\alpha$ -synucleinopathies (Mattson 2007). In PD pathology, it has been noted previously that neurons with higher levels of the calcium buffer, calbindin/D28K, are relatively spared (Yamada et al. 1990) and there may be an interplay between calcium homoeostasis and dopamine in mediating neuronal cell death (Mosharov et al. 2009). Indeed,  $\alpha$ -synuclein has been found to perturb cellular calcium homoeostasis (Hettiarachchi et al. 2009; Danzer et al. 2007). Recently, it was demonstrated that increased intracellular free calcium causes the formation of microscopically visible  $\alpha$ -synuclein aggregates in the cytoplasm of  $\alpha$ -syn-eGFP-transfected human glioma cells (Nath et al. 2011) and more recently that oxidative stress and raised calcium can act synergistically to cause  $\alpha$ -synuclein aggregation (Goodwin et al. 2012).

# 6 Ubiquitin and Ubiquitin Homologues

Ubiquitin, a small protein that attaches covalently to other proteins to target them to cellular pathways, such as proteasomal degradation, has long been known as

a major component of GCIs, and its presence has been linked both to the ubiquitin proteasome system, autophagy, and aggresome biogenesis. Recent studies indicate that polyubiquitin chains can be linked one ubiquitin to the next via many different ubiquitin functional groups, with each different type of ubiquitin chain conferring a different function on the modified protein carrying it, with the lysine-63-linked polyubiquitin targeting proteins to the aggresome and K48-linked ubiquitin chains targeting the proteasome (Kulathu and Komander 2012; Trempe 2011). Deubiquitinated  $\alpha$ -synuclein is targeted for autophagic degradation, while monoubiquitination preferentially targets  $\alpha$ -synuclein to the proteasome (Rott et al. 2011). In MSA, ubiquitin modifications of various proteins have been reported, including parkin and  $\alpha$ -synuclein, and compromise of the ubiquitin proteasome system has also been inferred based on reduced 20S proteasome subunit expression (Bukhatwa et al. 2010). Soluble aggregated forms of  $\alpha$ -synuclein have been shown to bind to the 19S cap of the 26S proteasome and inhibit proteasomemediated protein degradation (Snyder et al. 2003; Lindersson et al. 2004). In a recent study using a transgenic mouse model of MSA expressing  $\alpha$ -synuclein under an oligodendrocyte-specific promoter, Stefanova and co-workers (2012) have shown that proteasome inhibition caused by administration of a chemical inhibitor led to an MSA phenotype with pathological features, including fibrillar glial  $\alpha$ -synuclein aggregates, demyelination, and axonal disturbances, that closely mirror those of MSA.

# 6.1 SUMO-1

In 2005, it was reported that minute subdomains exist in GCIs that are marked by small ubiquitin-like modifier, SUMO-1 (Pountney et al. 2005c). SUMO (small ubiquitin-like modifier) is similar to ubiquitin in structure but with more diverse physiological functions. SUMOylation requires the coordinated actions of E1 (SUMO activating), E2 (SUMO-specific conjugating, Ubc9) and, in most cases, E3 (SUMO ligase, e.g., PIAS3) enzymes, and despite being a covalent protein modification, SUMOylation is readily reversed by SUMO-specific deSUMOylation enzymes, SENPs. Thus, the balance between Ubc9-mediated conjugation and SENP-mediated deconjugation determines the SUMOylation state (Martin et al. 2007). SUMOylation can inhibit protein binding or, alternatively, SUMOylation of target proteins can promote interactions with other proteins via SUMO interaction motifs (SIMs). SUMOylation functions to regulate nucleocytoplasmic transport by modification of RanGAP1 and subsequent association with the nuclear pore complex and also has roles in mitochondrial fission. SUMOylation modifies the mitotic spindle and, by mediating non-covalent interactions with dynamin, influences endocytosis. SUMO-1 has also been linked to a range of neuropathological protein species; indeed, both  $\alpha$ -synuclein and tau have been shown to be capable of SUMO-1 modification (Krumova and Weishaupt 2012). SUMO-1 was first identified in pathological inclusion bodies in neuronal intranuclear inclusions in neuronal

intranuclear inclusion disease, spinocerebellar ataxia type III, and Huntington's disease (Pountney et al. 2003). Recent studies have shown that neuron-specific expression of the SUMO-2 isoform can result in some SUMOylation of  $\alpha$ -synuclein and that the SUMOylated protein is less prone to aggregation (Krumova et al. 2011). However, the significance of this finding to glial cells is unclear. Several target proteins of SUMO-1 have been determined in intranuclear inclusions that could be associated with lysosomal or secretory pathways (Hsp90, Munc18, NSF, and dynamin-1) (Pountney et al. 2008), and it has been shown that SUMO-1 marks a subset of lysosomes in MSA and in glial cell culture models of proteolytic stress and protein aggregation. This tagging of lysosomes by SUMO-1 is correlated with the presence of cytoplasmic inclusion bodies and associated with Hsp90, suggesting SUMO-dependent modulation of the autophagy pathway in MSA (Wong et al. 2013).

## 7 Autophagy

Autophagy is an evolutionarily conserved bulk degradation process that involves lysosomes. There are three types of autophagy: microautophagy which is the direct uptake of cytosolic proteins or organelles into the lysosome, chaperone-mediated autophagy which involves the chaperone conjugate Hsc70 directing protein substrates to the lysosome, and also macroautophagy (autophagy) which involves the formation of autophagosomes that then fuse with endosomes or lysosomes (Klionsky et al. 2011). Significant progress has been made in delineating the molecular mechanisms of chaperone-mediated autophagy that includes targeting of proteins via  $EF1\alpha$  and interaction with GFAP at the lysosomal membrane (Kaushik et al. 2011). Recent studies have linked hsc70 also to a selective variant of autophagy that involves translocation of proteins via the multivesicular body (Sahu et al. 2011). Experiments have shown that autophagy is highly important in neural cells and disruption of the process can lead to aggregate formation and neurodegeneration, further highlighting its importance in neural tissue. The autophagy pathway has been implicated in neurodegenerative disease, and evidence of the pathways' involvement in MSA is seen in the presence of LC3 and lysosome components in glial inclusion bodies (Schwarz et al. 2012). Autophagy has been shown to contribute to aggresome clearance in neurodegenerative models and is believed to be an important cellular response to increased aggregating protein stress. Mouse knockouts of autophagy genes show progressive neurodegeneration, and disease-linked  $\alpha$ -synuclein forms have been shown to inhibit chaperonemediated autophagy (Rami 2009). Macroautophagy can enable the clearance of  $\alpha$ -synuclein aggregates, and disruption of the process can lead to aggregate formation and neurodegeneration, further highlighting its importance in neural tissue (Crews et al. 2010). All of these studies indicate a neuroprotective role for the autophagy pathway in neurodegenerative disease and raise the possibility of modulating this pathway as a therapeutic treatment.

#### 8 Oxidative Stress

Oxidative stress arises from an imbalance between the cellular processes that can generate powerful oxidants, such as superoxide and peroxynitrite, and the cellular defenses, such as glutathione, that normally combat them. Oxidative stress of neural cells has been implicated in the pathology of many neurodegenerative diseases, including MSA, and may play an important role in the development of MSA pathology, with the occurrence in GCIs of oxidized components, such as nitrated  $\alpha$ -synuclein and protein-bound 4-hydroxy-2-nonenal (Gómez-Tortosa et al. 2002; Norris and Giasson 2005; Shibata et al. 2010). Oxidative nucleic acid damage has also been reported in MSA (Kikuchi et al. 2002). Furthermore, oxidation is among the many different exogenous factors, including metal ions and lipids, that have been shown to induce the formation of potentially cytotoxic soluble  $\alpha$ -synuclein oligomers in cell culture and animal models of MSA (Brevdo et al. 2012; Stefanova et al. 2005, 2007; Riedel et al. 2007; Ubhi et al. 2009). Recently, the interaction between oxidation/oxidative stress and raised calcium was investigated, and it was found that these two factors act synergistically to induce  $\alpha$ -synuclein aggregates in vitro and in glial cells (Goodwin et al. 2012).

## 9 Conclusion

In conclusion, there is a strong burden of evidence pointing to a pivotal role of  $\alpha$ -synuclein aggregation in MSA development. Although it is likely that aggregated  $\alpha$ -synuclein forms occur at an early stage in the disease progression, there are many factors that have been identified, including oxidative stress and calcium dysregulation, that may participate in triggering the production of the pathogenic forms of the protein. The role of  $\alpha$ -synuclein posttranslational modifications, such as Ser-129 phosphorylation, is also unclear, especially with respect to the timing of the modifications either as a precursor to or a consequence of aggregate formation. Accumulation in GCIs of  $\alpha$ -synuclein and another small, acidic protein, metallothionein-III, even though mature oligodendrocytes do not normally express either protein, suggests that abnormal uptake of proteins by oligodendrocytes may be a feature of the etiology of MSA. Indeed, extracellular roles for both  $\alpha$ -synuclein and MT-III have recently been reported. Future studies will need to address the causes of (and potential remedies for)  $\alpha$ -synuclein aggregation in MSA as well as the origin of the protein in glial cells.

Another important aspect of MSA pathogenesis that will need to be probed further is the relationship between the formation of GCIs primarily in oligodendrocyte cells and the activation of neighboring astrocytes and microglia. Although it is clear that neuronal degeneration in the disease occurs as a downstream consequence of the glial cell disturbances, the relative importance of oligodendroglial dysfunction and demyelination compared to the influence of the inflammatory cytokines and astrocyte and microglial activation is unknown. Indeed, recent studies by Sofroniew and co-workers suggest that reactive astrogliosis may even have some neuroprotective as well as deleterious effects on neuronal survival in injured brain tissue (Sofroniew 2009; Verkhratsky et al. 2012).

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# Neurotoxic Vulnerability Underlying Schizophrenia Spectrum Disorders

Trevor Archer, Ulla Karilampi, Serafino Ricci, and Max Rapp-Ricciardi

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#### Abstract

Neurotoxic vulnerability that putatively contributes to the etiopathogenesis of schizophrenia spectrum disorders encompasses perinatal adversity, genetic linkage, epigenetic disadvantage, and neurodegenerative propensities that affect both symptom domains, positive, negative, and cognitive, and biomarkers of the disorder. Molecular and cellular apoptosis/excitotoxicity that culminates in regional brain loss, reductions in reelin expression, trophic disruption, perinatal adversity, glycogen synthase kinase-3 dysregulation, and various instances of

U. Karilampi

#### S. Ricci

Department of Anatomy, Histology, Forensic Medicine and Orthopaedics, Sapienza University of Rome, Rome, Italy

e-mail: serafino.ricci@uniroma1.it

T. Archer (🖂) • M. Rapp-Ricciardi

Department of Psychology, University of Gothenburg, Gothenburg, Sweden e-mail: Trevor.archer@psy.gu.se; max.rapp.ricciardi@psy.gu.se

Department of Psychiatry/Psychosis, Sahlgrenska University Hospital, Gothenburg, Sweden e-mail: ulla.karilampi@vgregion.se

oxidative stress all influence the final end point disorder. The existence of prodromal psychotic phases, structural–functional aspects of regional neuroimaging, dopamine signal overexpression, and psychosis propensity provide substance for neurodegenerative influences. The pathophysiology of schizophrenia spectrum disorder encompasses the destruction of normal functioning of the neurotrophins, in particular brain-derived neurotrophic factor (BDNF), dyskinesia of necessary movements, and metabolic–metabolomic and proteomic markers. Neurotoxic accidents combined with genetic susceptibility appear to play a role in interfering with normal neurodevelopment or in tissue-destructive neurodegeneration or both, thereby elevating the eventual risk for disorder tendencies and eventual expression.

Keywords

GFK • Imaging • Infection • MAM • Perinatal • Reelin • Trophic factors • Vulnerability

#### 1 Introduction

The contributions of neurotoxic circumstances to the development, predisposition, precipitation, and eventual expression of schizophrenia spectrum disorders may be sought at different levels of etiopathophysiogenesis. Several lines of evidence imply that these disorders originate from the abnormal development of the central nervous system (CNS) peculiar to the human condition, as exemplified by the entorhinal cortex that is involved in widespread cognitive networks (Akil and Lewis 1997; Arnold et al. 1991, 1997; Falkai et al. 2003). These may be illustrated in several respects. Both structural and functional abnormalities are associated with development of schizophrenia spectrum disorders: for example, lateral ventricular and third and fourth ventricular enlargements accompanied by reduced prefrontal, thalamic, and hippocampal volume have been observed consistently in the brain of patients afflicted by the disorder (Welch et al. 2011). These deficiencies include smaller-sized hippocampus and entorhinal cortex both associated with reduced neuronal body sizes (Baiano et al. 2008; Jonsson et al. 1999; Turetsky et al. 2003; Zaidel et al. 1997). Alexander-Bloch et al. (2010) have observed disrupted modularity and local connectivity of brain functional networks in childhood-onset schizophrenia. It has been shown too that both neuropsychologically near normal and neuropsychologically impaired patients had markedly smaller gray matter and larger third ventricle volumes than healthy comparison subjects. However, only neuropsychologically impaired patients had significantly less white matter and larger lateral ventricle volumes than healthy comparison subjects (Wexler et al. 2009). Thoma et al. (2009) showed that neuropsychological-functioning schizophrenic patients were impaired overall, and anterior hippocampus volume was smaller in the patient group. In healthy controls, the hippocampus-memory relationships involved the anterior and posterior hippocampus wherein correlations were significant for verbal memory measures. Among the schizophrenia patients, positive correlations were constrained to the posterior hippocampus. Negative correlations emerged between anterior hippocampus and verbal and visual memory measures. For both groups, cortical volume negatively correlated with age, but a negative correlation between age and hippocampus volume was found only among the schizophrenia patients. In their sample of adults with schizophrenia, atypical relationships between regional hippocampus volumes and episodic memory ability were found, as was an atypical negative association between hippocampus volume and age. Taken together, these diverse observations on the abnormal structural–functional relationships in schizophrenia spectrum disorders prompt notions of neurodegenerative disruptions during the early and later life cycles of brain development.

Imaging studies (positron emission tomography [PET]) or single photon emission tomography (SPET) and magnetic resonance imaging (MRI) attest to regional glutamatergic abnormalities in psychosis and are beginning to describe both the neurodevelopment of these abnormalities over the course of the illness and their response to therapeutic intervention (Egerton and Stone 2012). Fitzgerald (2012) discussed an N-methyl-D-aspartate Receptor (NMDAR) "suppression" hypothesis that focuses upon several aspects of evidence for NMDA influence in disorder: (1) epilepsy studies, (2) neurotoxicity studies, (3) mouse knockout studies of particular receptor subunits, (4) electrophysiological studies, (5) 2-deoxyglucose studies, and (6) functional brain imaging studies. Further, the activation of glycogen synthase kinase-3 (GSK-3) has been implicated in the cellular neurotoxicity of NMDAR antagonists (Large et al. 2011), with GSK-3 also implicated in schizophrenia (Brambrink et al. 2012; Pollard et al. 2012). Chan et al. (2012) have shown the critical role of glycogen synthase kinase-3 (GSK-3) in the expression of neurobehavioral aberrations associated with NMDAR hypofunction. They have suggested that GSK-3 inhibitors may ameliorate certain behavioral and cognitive dysfunctions in patients presenting schizophrenia. Neurotoxicity may be initiated by inflammatory agents: as described by Burd et al. (2012), infection-induced maternal immune activation leads to a fetal inflammatory response mediated through cytokines and are implicated in the pathophysiology of not only periventricular leukomalacia and cerebral palsy but also a spectrum of neurodevelopmental disorders, including attention-deficit hyperactivity disorder (ADHD), autism, and schizophrenia. Alternatively, Marona-Lewicka et al. (2011) have proposed an LSD model of schizophrenia whereby the chronic treatment of rats with low doses of LSD can serve as a new animal model of psychosis that may mimic the development and progression of the disorder.

Neurotoxic influences in schizophrenia spectrum disorders, to a large extent, would appear to be based upon circumstantial evidence rather than hard neurodegenerative loss of essential tissue, thereby rendering the associations somewhat tenuous. Notwithstanding these obvious constraints, there exists a sufficiency of observations from laboratory animal model studies, neuroimaging studies, and postmortem assays together with a neurodevelopmental viewpoint of disorder etiopathogenesis, suggesting that neurodegenerative processes contribute to height-ened predisposition and susceptibility to the precipitation of symptom domains through a staging notion (Archer et al. 2010a, b). Genetic risk factors may impact upon or disrupt developmental trajectories (Dauvermann et al. 2012; Sullivan et al. 2003). Indeed, genetic predispositions for neurodegenerative influences may be quite likely, since despite the absence of astrogliosis, impaired expression of astrocyte- and oligodendrocyte-related genes was observed in patients with major psychiatric disorders, including schizophrenia and mood disorders (Steiner et al. 2011). There is much support for the notion that schizophrenia spectrum disorders have their origin during gestation and/or in early infancy (Graff et al. 2011; Sawa and Snyder 2002). Insults during the early infancy or perinatal period, or even during the course of early-to-late childhood and adolescence, are implicated in the etiopathogenesis of the marked structural and functional abnormalities that distinguish the symptoms of disorder (Cannon et al. 2002a, b; Fontes et al. 2011; Geddes et al. 1999). The eventual contribution of neurodegenerative processes affecting the disorder must take into account the plethora of interactive factors and agents that include inherited and acquired attributes (Elia et al. 2011; Jablonka and Raz 2009). The disorder pathophysiology may involve the influences of several contributory processes, including cellular apoptosis/excitotoxicity, viral and bacterial infections, anoxic birth injury, maternal starvation, and other conditions exerting detrimental effects on neurogenesis. Both neurogenetic and epigenetic forces arising from geneenvironment interactions affecting developmental trajectories impart levels of predisposing vulnerability that may or may not lead to the manifestation of schizophrenic symptoms (Gratacos et al. 2007; Hoenicka et al. 2010) or those of related conditions (Bergman et al. 2011; Jacob et al. 2010). Whether or not the neurodegenerative aspect of the disorder may be neurogenetic or epigenetic, there does exist compelling evidence for expressions of the disruptive influence affecting normal development.

# 2 Regional Loss

At first sight, the accumulated evidence for progressive volumetric loss, e.g., in cortical volume/thickness (Honea et al. 2008; Goldman et al. 2009) and loss of functional connectivity (Meyer-Lindenberg et al. 2005; Yu et al. 2011), argues for the aspect of neurodegeneration in schizophrenia. Yet the possible impact of peristaltic and environmental agents must be understood prior to deriving a reliable conclusion (Meyer-Lindenberg 2011). For example, Goto et al. (2010) compared 18 patients (9 males, 9 females; age range: 13-52 years) with 18 healthy volunteers (9 males, 9 females; age range: 15-49 years) with no current or past psychiatric history, using magnetic resonance spectroscopy (MRS). They found that levels of N-acetylaspartate/Cr in the left basal ganglia and parieto-occipital lobe, but not in the frontal lobe, were significantly lower in patients with firstepisode schizophrenia psychosis than in control subjects. No differences were observed between the serum brain-derived neurotrophic factor levels of patients with first-episode schizophrenia psychosis and the healthy control subjects. The plasma levels of catecholamine metabolites, plasma 3-methoxy-4hydroxyphenylglycol (MHPG), but not homovanillic acid (HVA), were significantly lower in the patients with first-episode psychosis than in control subjects. In addition, a significantly positive correlation was observed between the levels of *N*-acetylaspartate/Cr of the left basal ganglia and plasma MHPG in all subjects. The authors concluded that brain *N*-acetylaspartate levels in the left basal ganglia and plasma MHPG levels were significantly reduced at the first episode of schizophrenia psychosis, indicating that neurodegeneration via noradrenergic neurons might be associated with the initial progression of the disease.

Regional loss of glutamate and glutamine in schizophrenic patients (Tayoshi et al. 2009) may offer evidence of an excitotoxic process in the disorder pathophysiology. The loss of parvalbumin-containing cells, an observation that has been reported consistently in postmortem analysis of the brains of schizophrenic patients following NMDAR blockade, offers support for this notion (Adell et al. 2012; Dean et al. 2011). Aoyama et al. (2011) have demonstrated that the loss of thalamic glutamate in the brain of schizophrenia patients correlated significantly with loss of gray matter in the middle and inferior frontal gyrus and temporal pole. Thalamic gray matter and glutamate losses were observed in individuals at risk for developing schizophrenia (Stone et al. 2009). Glutamate-glutamine ratios in the frontal cortices of adolescents at risk for the disorder are correlated, as well as performance scores on the Global Assessment of Functioning Scale (Tibbo et al. 2004). It has been suggested that the elevated glutamine levels in never-treated patients followed by the decreased level of thalamic glutamine and gray matter loss in connected regions may be indicative of either neurodegeneration or a plastic response to reduced subcortical activity (Theberge et al. 2007). There appears to be a consensus of sorts for an excitotoxic contribution primarily in early stage schizophrenia (Bustillo et al. 2010). Glutamatergic alterations and loss of cortical gray matter originating from basal ganglia-corticothalamic circuits may be the expressions of excitotoxicity through these regions. First-episode schizophrenic patients have exhibited increased membrane breakdown in magnetic resonance spectroscopy applying labeled phosphorous (Miller et al. 2009).

#### 3 Perinatal Adversity

Several perinatal insults have been shown to induce schizophrenia-like symptoms in a laboratory setting (Archer 2010). Bilateral excitotoxin-induced lesions, administering ibotenic acid to the ventral hippocampus typically to 7-day-old rat pups, induces disruptions of prepulse inhibition (PPI) to the acoustic startle response, a model of sensory gating in rodents, hyper-responsiveness to stressful stimuli, supersensitivity to dopamine (DA) agonists, and NMDAR antagonists (Lipska et al. 1992, 1995, 2001; Tseng et al. 2007, 2009). Additionally, persistent deficits in working memory and spatial navigational learning (O'Donnell et al. 2002), together with hyper-responsiveness to stimulation of the ventral tegmental area (Lillrank et al. 1999; Lipska and Weinberger 1994), have all been observed in neonatal ventral hippocampus-lesioned rats. Prenatal administration of methylazoxymethanol (MAM) disrupts early brain development with structural and functional

abnormalities in the cortex and hippocampus, many of which bear similarities to schizophrenia, expressed in the adult rats (Flagstad et al. 2005; Lodge and Grace 2008; Mohammed et al. 1986a, b; Moore et al. 2006). MAM induces programmed cell death in neuroepithelial-proliferating cells (Cattabeni and Di Luca 1997), without causing any teratogenic effects in any other organs (Balduini et al. 1991). It is generally administered on embryonic, gestation, day 17 (E17) as this is when neurogenesis peaks in the hippocampal formation (Bayer and Altman 2004), and the time when projections between the hippocampus and entorhinal cortex are established (Ceranik et al. 1999). Hippocampal formation disruption is central to schizophrenia pathophysiology (Harrison 2004). The utility of prenatal MAM as an animal model of the disorder highlights the neurodegenerative aspect (Chen and Hillman 1986; Ferguson and Holson 1997; Ferguson et al. 1996; Virgili et al. 1997), particularly with regard to the neurocognitive deficits (Fiore et al. 2002; Gourevitch et al. 2004: Lee and Rabe 1992; Leng et al. 2005). The model has been validated by several studies (cf. Gill et al. 2011; Hradetzky et al. 2011; Lodge and Grace 2012; Valenti et al. 2011). Neurophysiological studies confirm that MAM-treated rats show an augmented DA system function consistent with enhanced DA signaling underlying positive symptoms of disorder and defective hippocampal interneurons (Konradi et al. 2011; Lahti et al. 2006). Similar to patients presenting cognitive deficit profiles in schizophrenia spectrum disorder (Karilampi et al. 2007, 2011), MAM induces marked deficits on an attentional-shift task (analogous to the Wisconsin Card Sorting Task) and on a differential reinforcement of low rate of responding (DRL-20 performance, analogous to continuous performance) task (Featherstone et al. 2007). Prenatal MAM administration disrupts early cortical development causing deficits in medial prefrontal functions at adult ages (Goto and Grace 2006). Sanderson et al. (2012) obtained a reduction in synaptic innervation and synaptic transmission in the dorsal hippocampus of MAM (E17)-treated rats, accompanied by a pronounced increase in CA1 pyramidal neuron excitability. These effects of the prenatally administered neurotoxin interrupt normal neuronal development, stress and immune response systems, and signal transduction mechanisms (Ciani et al. 2003; Ferrer et al. 1997; Lafarga et al. 1997; Rice and Barone 2000), thereby reinforcing its utility as a model of schizophrenia (Gill et al. 2011; Le Pen et al. 2011) and implying the neurotoxic aspect of the disorder.

Traumatic/adverse experiences during infancy and early childhood compromise healthy brain development in several domains, including cognitive, emotional, and motor. Traumatic environmental confrontation during early neurodevelopment, such as parental loss (maternal separation), that induces a marked provocation of the hypothalamic–pituitary–adrenal (HPA) axis was found to cause schizophrenialike deficits, e.g., impairments of sensory gating, symptoms in the rats as adults (Ellenbroek et al. 1998), as well as other expression of the disorder (Furukawa et al. 1998). Rosenberg et al. (2007) surveyed 569 adults presenting schizophrenia with regard to adverse childhood events (including physical abuse, sexual abuse, parental mental illnesses, loss of a parent, parental separation or divorce, witnessing domestic violence, and foster or kinship care), evaluating the relationships between cumulative exposure to these events and psychiatric, physical, and functional outcomes. They found that increased exposure to adverse childhood events was strongly related to psychiatric problems in both cognitive and emotional domains (suicidal thinking, hospitalizations, distress, and posttraumatic stress disorder), substance abuse, physical health problems (HIV infection), medical service utilization (physician visits), and poor social and daily care functioning (homelessness or criminal justice involvement). Similarly, Morgan et al. (2007) observed that separation from, and/or the death of, one parent prior to the age of 16 years was strongly associated with a twofold to threefold increases in the risk of psychosis. It was shown that the strength of these associations was similar for White British and Black Caribbean (but not Black African) subjects, although separation from (but not death of) a parent was more common among Black Caribbean controls than White British controls. Finally, a 10-year cohort of high security hospital patients who had either a personality disorder or schizophrenia found a rate of child-parent separation of 178/289 (62 %) in the schizophrenia group compared with a rate of 119/147 (81 %) in the personality disorder group of patients that had been separated from one or both parents before the age of 16 (Pert et al. 2004). Several studies have described the extreme adverse effects of perinatal starvation/malnutrition involving severe deficits of necessary nutrients, including folate, essential fatty acids, retinoids, vitamin D, and iron, the "nutritional hypothesis of schizophrenia" (Brown and Susser 2008). Additionally, elevated prenatal homocysteine level provides a plausible risk factor for schizophrenia because of its partial antagonism of N-methyl-p-aspartate receptors under physiologic glycine concentrations and its association with abnormal placental function and pregnancy complications (Bleich et al. 2007; Brown and Susser 2005; Brown et al. 2007).

An abundance of clinical and preclinical (animal modeling) findings posit the notion that immune system dysfunction(s) – related to genetic, environmental, and neurobiological influences – may play a role in the etiology of schizophrenia in a subset of patients (Richard and Brahm 2012). Exposure to prenatal immune challenge induces early pre- and postnatal alterations in peripheral and central inflammatory response systems. Consequentially, these alterations may disrupt the normal development and maturation of neuronal systems from juvenile to adult stages of life, thereby affecting brain structure and function. Developmental neuroinflammatory processes of this type may adversely affect processes that are pivotal for normal brain maturation, including myelination, synaptic pruning, and neuronal remodeling, all of which occur to a great extent during postnatal brain maturation (Meyer 2011). Brown and co-workers (Brown and Derkits 2010; Brown and Patterson 2011; Brown et al. 2009, 2011) and others (Arias et al. 2012; Nielsen et al. 2011; Tedla et al. 2011) have conducted birth cohort studies to address whether in utero exposure to infectious agents, prospectively documented by biomarker assays of archived maternal sera and by detailed obstetric records, confers an increased risk of schizophrenia in adult offspring. Prenatal exposure to influenza, elevated toxoplasma antibody, rubella, genital/reproductive infections, and other infections have been associated with an increased risk of schizophrenia among offspring. Animal models have supported these epidemiologic findings by revealing that maternal immune activation causes phenotypes analogous to those found in patients with schizophrenia (Brown 2011). In a population-based study, Benros et al. (2011) have shown that autoimmune disease conditions and the number of infections requiring hospitalization pose risk factors for schizophrenia. Increased risk for the disorder was compatible with an immunological hypothesis in subgroups of schizophrenia patients.

# 4 Reelin Reduction

Reelin, present throughout the brain, spinal cord, blood, and other organs and tissues, is a large secreted extracellular matrix glycoprotein that facilitates regulation of processes involving neuronal migration and positioning in the developing brain by controlling cell-cell interactions and maintains its role in the adult brain. Reelin modulates synaptic plasticity and stimulates dendrite and dendritic spine development and regulates the continuing migration of neuroblasts generated at sites of adult neurogenesis (Niu et al. 2008). Expression of the protein is found to be significantly lower in schizophrenia and psychotic bipolar disorder (Maloku et al. 2010; Pisanté et al. 2009; Suzuki et al. 2008). Several studies have shown that in schizophrenia spectrum disorders, reelin expression in several brain regions, such as the cerebral cortex and hippocampus, is reduced (Eastwood and Harrison 2006; Fatemi et al. 2000; Impagnatiello et al. 1998). Reelin expression is reduced in the prefrontal cortex (PFC), hippocampus, caudate nucleus, and cerebellum of patients presenting psychotic illness (Guidotti et al. 2000; Tueting et al. 2006); GABAergic interneurons express reelin in several brain regions (Miyata et al. 2010; Pascual et al. 2004; Pesold et al. 1998). In order to induce prenatal infection (gestation days 15 and 16) in rat offspring, Nouel et al. (2012) exposed them to lipopolysaccharide, a bacterial endotoxin, and then assessed the number of reelin-immunoreactive (reelin+) cells and glutamic acid decarboxylase 67 (GAD67) immunoreactive (GAD67+) cells that were present in the hippocampus at postnatal days 14 and 28. They observed decreases in both reelin+ and GAD67+ expression in the dentate gyrus of infected animals at postnatal day 14 and concluded that prenatal infection caused postnatal reductions in postnatal reelin and GAD67. Prenatal exposure to influenza, elevated toxoplasma antibody, genital/reproductive infections, rubella, and other pathogens have been shown to be associated with schizophrenia (Brown 2012; Ducharme et al. 2012; Kneeland and Fatemi 2012; Macêdo et al. 2012).

Reelin regulates radial neuronal migration in the embryonic brain, promotes dendrite outgrowth in the developing postnatal forebrain, and strengthens synaptic transmission in the adult brain. In contrast, reelin deficiency induces specific defects in the molecular composition of the synapses in the adult brain (Ventruti et al. 2011). The MAM E17 model of schizophrenia was observed to cause whole neuronal density changes from the PFC and parvalbumin-positive-cell density changes from the hippocampus, and similar deficits were seen in the PFC of schizophrenic patients (Lodge et al. 2009; Moore et al. 2006). Matricon et al. (2010) exposed rat fetuses to MAM during E17 and then studied behavioral parameters and biomarkers in the adult animal. They found that behavioral

characteristics and neuropathology pertaining to the PFC similar to clinical features of the disorder were in evidence. In comparison with untreated offspring, MAM-exposed offspring showed a reduction in the volume of entorhinal cortex, hippocampus, and mediodorsal thalamus that was associated with reduced neuronal soma. The entorhinal cortex also showed laminar disorganization and neuronal clusters. Reelin methylation in the hippocampus formation was decreased although reelin + neurons and reelin expression were unchanged. They concluded that MAM-treated animals have a diminished neuropil possibly arising from abnormal neurite formation. The deterioration of the neuropil in cortical regions is implicated in the neuropathology of schizophrenia (Somenarain and Jones 2010). Protein loss in the disorder induces dendrites and spines to deteriorate in the dorsolateral PFC that is involved in several key cognitive domains.

# 5 Trophic Disruption

Brain-derived neurotrophic factor (BDNF) is essential for the maintenance of functional neurons, regulating growth, differentiation, synaptic connectivity, neurorepair, and longevity (Altar et al. 1997; Niu and Yip 2011). BDNF, central for the survival and differentiation of midbrain dopaminergic neurons (Hyman et al. 1991) and phenotypic differentiation of locus coeruleus noradrenergic neurons (Traver et al. 2006), has emerged as an important biomarker for schizophrenia spectrum disorder (Pae et al. 2012; Favalli et al. 2012; Zhang et al. 2012b). Sixtythree patients presenting schizophrenia symptoms were compared with 52 age- and sex-matched healthy controls for performance on a battery of neuropsychological tests by Niitsu et al. (2011). They found that although there were no significant differences in serum BDNF levels between normal controls and schizophrenic patients, serum BDNF levels for normal controls, but not schizophrenic patients, showed negative correlations with verbal working memory. On the other hand, serum BDNF levels of schizophrenic patients indicated positive correlations with the scores of the Scale for the Assessment of Negative Symptoms (SANS) and the Information subtest scores of Wechsler Adult Intelligence Scale Revised (WAIS-R). Serum BDNF levels were related with the impairment of verbal working memory and negative symptoms in patients with schizophrenia. In 22 acute schizophrenic patients and 22 age-matched healthy volunteers, Lee et al. (2011) observed significantly reduced serum levels in the unmedicated schizophrenic patients (n = 22; 4.38  $\pm$ 2.1 ng/mL) compared to the age-matched healthy volunteers. The percentage change of BDNF (increase, 173  $\% \pm 110$ ) correlated negatively with the percentage change of PANSS score with BDNF increase during psychotic treatment, as shown by several others (e.g., Pedrini et al. 2011; Yoshimura et al. 2007, 2010). Reduced BDNF concentrations have been reported in chronic antipsychotic-treated patients (Grillo et al. 2007; Ikeda et al. 2008; Rizos et al. 2010), neuroleptic-free (Palomino et al. 2006) and neuroleptic-naïve (Chen and Huang 2011; Rizos et al. 2008) patients. Finally, the levels of peripheral BDNF seem to be altered in first-episode patients presenting psychosis as well as in chronic schizophrenic patients. Although a paucity

of studies have reported changes in peripheral BDNF levels following antipsychotic treatment, the role of Val66Met polymorphism in BDNF has been found to contribute to structural and functional plasticity in schizophrenia (Buckley et al. 2007, 2011).

Long-term administration of antipsychotic agents induces tardive dyskinesias (TDs), a syndrome composed of involuntary, hyperkinetic, and abnormal movements often expressed through excessive chewing or dancing/foot shuffling behaviors (Jafari et al. 2012; Correll and Schenk 2008), Lower serum concentrations of BDNF, inversely correlated with Abnormal Involuntary Movement Scale (AIMS) scores, have been observed in patients presenting TDs (Tan et al. 2005). Additionally, associations between TD and BDNF polymorphisms have accumulated (Park et al. 2009; Zai et al. 2009). Yang et al. (2011) compared serum BDNF levels in schizophrenic patients with (n = 129) and without (n = 235) TDs with healthy controls (n = 323), with concurrent assessment of AIMS and the Positive and Negative Symptoms Scale (PANSS). Schizophrenic patients presenting TDs showed lower BDNF concentrations, while schizophrenic patients lacking TDs had BDNF levels similar to that of healthy controls. Lower BDNF serum concentrations correlated with higher PANSS negative subscores, but not with AIMS scores. The authors concluded that reduced BDNF concentrations may be associated with greater TD pathophysiology and negative symptoms in schizophrenia. Zhang et al. (2010) have indicated that schizophrenic patients with TD presented higher serum levels of \$100B, a calcium-binding protein, than normal, healthy controls and those patients without TD. The serum S100B levels were positively correlated with AIMS scores in patients with TD. The authors concluded from these results that increased S100B levels may be related to the neurodegenerative aspect of disorder, associated with TD pathophysiology. TDs are generally indicative of neurodegenerative pathologies (e.g., Miller and Chouinard 1993).

# 6 GSK Regulation

The glycogen synthase kinase-3 s (GSK,  $\alpha$ , and  $\beta$ ), involved in glycogen metabolism, are expressed ubiquitously over various tissues and are abundant in brain tissue (Jaworski et al. 2011; Perez-Costas et al. 2010). Kaidanovich-Beilin and Woodgett (2011) have discussed the fundamental roles for these protein kinases in memory, behavior, and neuronal fate determination. It has been argued that Schizophrenia-1 (DISC1) represents a candidate gene for neuropsychiatric disorders and an influential involvement during brain structural and functional development. Singh et al. (2011) have observed that DISC1 variants are implicated in the loss of function in Wnt/GSK-3 $\beta$  signaling and thereby interrupt the course of brain development. Common DISC1 polymorphisms (variants) are associated with neuropsychiatric phenotypes including altered cognition, brain structure, and function. The contribution of the DISC1 gene to neural development and the eventual neurodevelopment of schizophrenia risk has been studied (Wexler and Geschwind 2011). GSK-3, a regulator of a wide range of cellular processes, plays a key dual role in apoptosis with putative contribution to schizophrenia neuropathology (Emamian et al. 2004) and other neurodegenerative disorders (Avila et al. 2004; Berger et al. 2005; Onishi et al. 2011). Its role in the apoptotic signaling underlying excessive cell death may imply a neurodegenerative involvement in schizophrenia spectrum disorders, at least during the early stages of development (Gomez-Sintes et al. 2011). Astrocytic glycogenolysis and glycogen mobilization are implicated in normal brain function (Brown and Ransom 2007; Brown et al. 2003; Swanson 1992) and neuronal energy requirements (Brown et al. 2004; Wender et al. 2000), and not least in the necessities of cognitive function (Gibbs et al. 2006; Hertz et al. 2003; Suzuki et al. 2011). The links between glycogen metabolism and the glutathione (GSH) system have been studied (Dringen and Hirrlinger 2003): Shinohara et al. (2010) have shown that activation of GSK-3 $\beta$  is a key mediator of the initial phase of acetaminophen-induced liver injury through modulating glutamate cysteine ligase (GCL) and myeloid cell leukemia sequence-1 degradation. GSH deficits occur both in neurodegenerative conditions and in schizophrenia (Ballatori et al. 2009; Do et al. 2009; Dringen et al. 1997), and a GCL gene polymorphism, affecting GSH synthesis, is associated with the disorder (Tosic et al. 2006). In this regard, Lavoie et al. (2011) have found that glucose metabolism and glycogen utilization are dysregulated in astrocytes showing chronic deficit in GSH, implying dysfunctional brain energy metabolism in schizophrenia.

## 7 Oxidative Stress

Dysregulation in reactive oxygen species (ROS) and reactive nitrogen species (RNS) evidenced by lipid peroxidation in plasma, red blood cells, blood platelets, and cerebrospinal fluid have been observed in schizophrenic patients (Dietrich-Muszalska et al. 2005; Li et al. 2006; Yao et al. 1998, 2001). For example, Dietrich-Muszalska et al. (2005) found that oxidative stress occurred as reflected by superoxide dismutase (SOD) in platelets from schizophrenic patients; suppressed SOD activity was associated with enhanced ROS generation and lipid peroxidation as compared with healthy volunteers. In the acute phase of psychosis in schizophrenic patients, levels of isoprostanes were remarkably high compared with the healthy control group (Dietrich-Muszalska and Olas 2007). The isoprostanes are prostaglandin-like compounds formed in vivo from the free radical-catalyzed peroxidation of essential fatty acids (primarily arachidonic acid) without the direct action of cyclooxygenase (COX) enzyme, and increased levels are associated with oxidative stress and oxidative damage of lipids. During the same acute phase of psychosis, biomarker concentrations of oxidation/nitration proteins in plasma were markedly higher in patients, as too was the homocysteine (Hcys) level (Dietrich-Muszalska et al. 2009): Heys may function as an oxidant both in vitro and in vivo (Carluccio et al. 2007; Olas et al. 2008) since there is a reduced plasma status of antioxidants in schizophrenic patients (Dietrich-Muszalska and Kontek 2010). Significant positive relationships between brain atrophy and plasma Hcys levels have been obtained (Bleich et al. 2002; Sachdev et al. 2002), with Hcys metabolism implicated in schizophrenia pathophysiology (Song et al. 2007).

Furthermore, increased third trimester Hcys concentrations could well enhance the risk for development of the disorder through structural and functional alterations (Brown and Ransom 2007). In this regard, Dietrich-Muszalska et al. (2012) investigated, in schizophrenic patients, the effect of elevated Hcys on parameters of oxidative stress, including thiobarbituric acid reactive substances (TBARS), levels of carbonyl groups in plasma proteins, and levels of 3-nitrotyrosine in plasma proteins. They observed that in these patients, plasma Hcys was greater than in controls. Also, the biomarkers of oxidative/nitrative stress, carbonyl groups and 3-nitrotyrosine, were enhanced significantly.

The pathophysiology of schizophrenia spectrum disorders may be the result of deregulation of synaptic plasticity, with downstream alterations of inflammatory immune processes regulated by cytokines, impaired antioxidant defense, and increased lipid peroxidation. Pedrini et al. (2012) examined serum oxidative stress markers and inflammatory cytokines in early and late phases of chronic SZ. They studied patients at early stage (within first 10 years of a psychotic episode), 39 at late stage (minimum 10 years after diagnosis of SZ), and their respective matched controls. From each patient/subject 5 mL blood samples were collected (venipuncture) for assay of TBARS, total reactive antioxidant potential (TRAP), protein carbonyl content (PCC), Interleukins 6 and 10 (IL-6, IL-10), and tumor necrosis factor alpha (TNF-alpha). TBARS, IL-6, and PCC levels were significantly higher in schizophrenic patients at early and late stages than in healthy controls. No differences for TRAP and TNF-alpha levels in patients with SZ at early and late stages than in controls were obtained. IL-10 levels were decreased in patients at late stage and a decrease trend in early stage was found. Their findings offer evidence consistent with comparable biological markers across the chronic disorder. Other results suggest that oxidative stress may be involved in the pathophysiology of the disorder and associated functioning, e.g., cognitive impairment: Zhang et al. (2012a) found that in schizophrenic patients' plasma, total antioxidant status was inversely associated with some domains of cognitive deficits in schizophrenia, such as attention and immediate memory. It is known too that adult neurogenesis transiently generates oxidative stress which may further contribute to the pathophysiology (Walton et al. 2012). Finally, Sullivan and O'Donnell (2012) have advanced the notion of an excitation-inhibition imbalance hypothesis of the neurodevelopment of symptom profiles in schizophrenia spectrum disorders; they have examined ongoing efforts that investigate the determination of whether or not cortical inhibitory interneurons are affected by oxidative stress during brain development.

#### 8 Saitohin Gene Expression

Finally, Saitohin, an intronless gene nested within the human tau gene that contains a single nucleotide polymorphism (A/G), may be involved in the pathophysiology of neurodegenerative disorders (Combarros et al. 2003). The Saitohin gene is unique to humans and their closest relatives although function is not currently understood. It contains a single polymorphism (Q7R); the Q allele is human specific

and confers susceptibility to several neurodegenerative diseases (Gao et al. 2005; Wang et al. 2011). Certain neurodegenerative disorders, particularly Alzheimer's disease and progressive nuclear palsy, have been found to be associated with the presence of the Q allele of the Saitohin gene (Janković et al. 2008; Lorenzi et al. 2010; Schutte et al. 2011). In a sample of 48 schizophrenic patients and 47 healthy controls, Bosia et al. (2011) tested the role of saitohin polymorphism as a concurring factor of cognitive decline among these patients using the Wisconsin Card Sorting Test for executive functioning. They observed a significantly greater frequency of G allele among patients with frontotemporal dementia and schizophrenic patients presenting impaired Wisconsin Card Sorting Test performance. They concluded that the products expressed by the Saitohin gene affected core frontal executive function likely through mechanisms occurring through neurodevelopment with neurodegenerative outcome.

#### 9 Conclusion

The myelin-centered model of the human brain's extensive and effective myelination implies the availability of higher metabolic resources underlying evolutionary adaptations resulting in the quadratic (inverted U) myelination trajectory that is peaking during the sixth decade of life (Bartzokis 2011). The notion proposes further that optimal brain function depends on exquisite action potential synchronization that myelin makes possible and that myelin's exceptional vulnerability to subtle metabolic/oxidative abnormalities may promote both developmental and degenerative diseases. It remains a possibility that in the interventions for schizophrenia spectrum disorders, the widespread use of psychotropic treatments may induce underappreciated CNS metabolic and neurotransmitter effects on myelination, its plasticity, and neuronal repair (Bartzokis 2011).

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# **Neurotoxicity of Methamphetamine**

# Rosario Moratalla, Sara Ares-Santos, and Noelia Granado

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#### Abstract

Recreational consumption of the highly addictive psychostimulant methamphetamine is becoming a serious public health problem worldwide. Recent estimates indicate that methamphetamine abuse has increased in the last decade and that only cannabis is used by a greater number of consumers. Despite its popularity, methamphetamine is a known neurotoxin that damages dopaminergic terminals

R. Moratalla (🖂) • S. Ares-Santos • N. Granado

Instituto Cajal, Consejo Superior de Investigaciones Científicas (CSIC), and CIBERNED, ISCIII, Madrid, Spain

e-mail: moratalla@cajal.csic.es; saraares@cajal.csic.es; ngranado@cajal.csic.es

in the striatum, as indicated by reductions in striatal levels of dopamine and its metabolites and a sustained decrease in the expression of markers for dopaminergic terminals such as TH and DAT. In addition, methamphetamine affects the cell bodies of these same dopaminergic neurons in the substantia nigra, resulting in cell loss. The mechanisms underlying dopaminergic neurotoxicity are the focus of intense research, and knowledge in this area has expanded in recent decades. Evidence from previous studies points to dysregulation of dopamine, oxidative stress, DNA damage, and mitochondrial dysfunction as the main causes of methamphetamine neurotoxicity. The dopamine receptors D1 and D2 also play an important role in methamphetamine-induced neurotoxicity since inactivation of either receptor is neuroprotective against methamphetamine. Recent results from clinical research indicate that methamphetamine abusers have a higher risk of developing Parkinson's disease; this is in keeping with results in laboratory animals and confirms the persistence of methamphetamine-induced dopaminergic injury. These findings suggest that neuroprotective strategies that are effective against methamphetamine-induced toxicity are also promising candidates for preventive therapy for Parkinson's disease and other persistent dopaminergic injuries.

#### Keywords

Amphetamine derivatives • Designer drugs • Dopamine • Drug addiction • Methamphetamine • Neurotoxicity • Parkinson's disease • Psychomotor stimulants

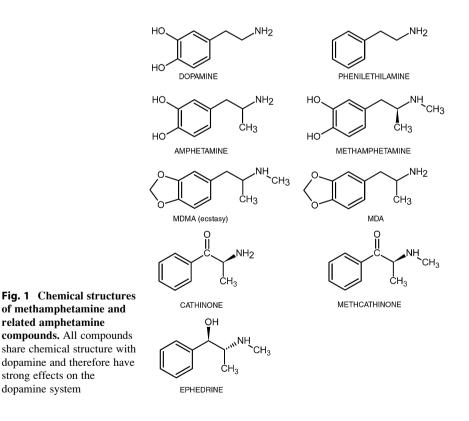
List of Abbreviations					
ATS	Amphetamine-type stimulants				
$D1R^{-/-}$	D1R knockout mice				
D1R	Dopamine D1 receptor				
$D2R^{-/-}$	D2R knockout mice				
D2R	Dopamine D2 receptor				
DAT	Dopamine transporter				
DOPAC	3,4-Dihydroxyphenylacetic acid				
Gclm	γ-Cysteine ligase modulatory subunit				
Gclc	γ-Cysteine ligase catalytic subunit				
GPx	Glutathione peroxidase				
HVA	Homovanillic acid				
MDMA	3,4-Methylendioxymethamphetamine also called "ecstasy"				
Nrf2	Nuclear factor-erythroid 2-related factor 2				
PD	Parkinson's disease				
ROS	Reactive oxygen species				
SNpc	Substantia nigra pars compacta				
TH	Tyrosine hydroxylase				
VMAT2	Vesicular monoamine transporter				

### 1 Introduction

## 1.1 Background, Medical Use, and Epidemiology

Methamphetamine (*N*-methyl-1-phenylpropan-2-amine) is a synthetic drug first used clinically at the beginning of the twentieth century. Therapeutic use of methamphetamine was based on its sympathicomimetic properties, but its current illicit use as a recreational drug in several countries around the world is due to its psychostimulant effects. It is also used for weight loss and for enhancing alertness, focus, motivation, and mental clarity for extended periods of time. Methamphetamine is structurally related to the neurotransmitters dopamine and phenylethylamine, and to other psychostimulant drugs like amphetamines (Fig. 1).

Methamphetamine has proven to be highly addictive and its abuse can result in severe psychological and physical dependence. It is therefore classified as a Schedule II drug under the Convention on Psychotropic Substances. Methamphetamine abuse is increasingly recognized as a major global health problem (Degenhardt et al. 2008). The 2012 World Drug Report (UNODC 2012) suggests that the global use of amphetamine-type stimulants (ATS; methamphetamine,



amphetamine, and methcathionine) (excluding "ecstasy") is second only to cannabis. They were used by 14–53 million people in 2010 equivalent to 0.3-1.2 % of the global population aged 15–64 years.

# 1.2 Administration Routes and Patterns of Methamphetamine Use

Methamphetamine comes in several forms. The hydrochloride salt of methamphetamine is a white, crystalline, bitter, odorless powder. It is water-soluble and strongly hygroscopic (absorbs water quickly). The common street names "speed," "Meth," or "chalk" refer to the salt, while "crystal," "crystal Meth," and "ice" refer to crystalline methamphetamine, a form purer than the powder. Methamphetamine is also known by a variety of other names, including shabu, batu, D-meth, tina, and glass. Methamphetamine freebase is oily and is uncommon on the street.

Methamphetamine can be taken orally (in pill form), by intravenous injection, smoking, snorting (in powder form), or by anal or vaginal insertion of a suppository. The effects experienced by the user last 6–8 h, depending on the rate at which methamphetamine reaches the blood, which depends on the route of administration. The faster the drug reaches the blood, the greater the "high" and other effects experienced by the user.

Following administration by any route, methamphetamine is distributed through most organs, including the lungs, liver, and stomach. Moderate levels reach the brain by crossing the blood-brain barrier. It also crosses the placenta and is secreted into breast milk. Methamphetamine is metabolized in the liver, with the main metabolites including the active compounds amphetamine, 4-hydrooxyamphetamine, and norephedrine. The concentration of amphetamine metabolite peaks at 10–24 h post-administration.

Methamphetamine abuse has two distinct use patterns. The first, characterized by low intensity use, does not confer psychological dependence. The second, known as "binge" use, consists of repeated redosing, usually by inhalation or injection, for several days in a row; generally, withdrawal symptoms occur when drug delivery is stopped abruptly.

### 1.3 Methamphetamine: Mechanism of Action and Effects

Methamphetamine's effects on the body are due to its structural resemblance to the neurotransmitter dopamine (Fig. 1). It easily crosses the blood–brain barrier, reaches the brain, and enters the axons of dopaminergic neurons through the dopamine transporter DAT and by passive diffusion (Granado et al. 2011a; Krasnova and Cadet 2009). Once inside the axon, methamphetamine triggers the release of dopamine from synaptic storage vesicles, resulting in an unusually high concentration of dopamine in the cytoplasm (Krasnova and Cadet 2009). Methamphetamine cannot directly activate dopamine receptors; rather, it acts as an indirect

dopamine agonist that increases dopamine concentration in the synapse by increasing dopamine release and blocking dopamine uptake. Methamphetamine also releases norepinephrine and serotonin by a similar mechanism; however, in the brain, methamphetamine is selectively concentrated in norepinephrine and dopamine nerve terminals because it is a substrate for the molecular transporters present in these terminals.

As a consequence of this increased release of dopamine in several areas of the brain, methamphetamine produces a number of acute psychological effects including euphoria (also known as "flash" or "rush", and lasting only several minutes). After this first short period, other feelings and behaviors may appear, including a false sense of self-confidence and power (delusions of grandeur), loquacity, moodiness, irritability, anxiousness, nervousness, aggressiveness, and violent behavior. Methamphetamine consumption has many acute adverse physical effects, including hyperthermia, increase in blood pressure and heart rate, mydriasis (pupil dilatation), logorrhoea, grinding teeth (trismus and bruxia), gastrointestinal irritation, appetite loss, itching, welts on skin, hyperactivity, involuntary body movements, irreversible damage to blood vessels in the brain resulting in cerebrovascular accidents, arrhythmia, tachycardia, cardiovascular collapse, and death. The most common symptoms of chronic methamphetamine abuse include temporomandibular joint syndrome, tooth erosion, and myofacial pain, all manifestations of acute trismus and bruxia. Other long-term symptoms are loss of appetite, weight loss, accelerated aging, nose bleeding, and "Meth mouth," an oral disease characterized by tooth erosion, extensive caries, decayed surfaces, missing teeth, tooth wear, plaque, and calculus (Fig. 2). Methamphetamine is highly addictive, and its use can result in tolerance: The effects decrease gradually with chronic use; thus, increased dosages are required to achieve the desired effects.

# 2 Methamphetamine Induces Neurotoxicity

Repeated methamphetamine administration results in neurotoxicity, primarily affecting dopaminergic neurons in the nigrostriatal system as reflected by longlasting reductions in levels of dopamine and its metabolites (DOPAC and HVA), dopaminergic markers such as tyrosine hydroxylase (TH) (the rate-limiting enzyme for dopamine synthesis), and DAT (Krasnova and Cadet 2009) (Fig. 3).

Several studies have demonstrated dopaminergic axon loss in the striatum after repeated methamphetamine use, indicated by loss of TH and DAT immunoreactivity. Although there is partial recovery of axonal TH and DAT immunoreactivity, some loss persists for long periods. Other amphetamine compounds such as ecstasy (MDMA) also produce this persistent axonal loss, which correlates with dopaminergic cell body loss in the substantia nigra pars compacta (SNpc), as demonstrated by rigorous stereology/cell counts with TH and Nissl staining and by use of cell death markers such as Fluorojade (Granado et al. 2008a). Apoptotic cell bodies, an irrefutable marker of cell death, have also been observed in the SNpc of methamphetamine-treated mice (Ares-Santos et al. 2012, 2013; Granado et al. 2011a).



**Fig. 2** Methamphetamine abuse can produce accelerated aging, "METH mouth," and "METH mites." (a). Physical aspect of a woman at several time points during 4 years of methamphetamine abuse. (b). Case of "Meth Mouth". (c). Sores on the skin known as "meth mites," which result from scratching to relieve the feeling of having small bugs under the skin that methamphetamine abusers may experience (Taken from drug enforcement administration www. dea.gov)

Interestingly, the compartments of the mouse striatum – the striosomes and matrix – differ in their vulnerabilities to methamphetamine, (Granado et al. 2010) (see Fig. 4). Striosomes, which are connected to the limbic system and are functionally associated with reward-related behaviors and emotional events (White and Hiroi 1998), are more vulnerable to methamphetamine-induced dopaminergic terminal loss than the matrix, which is connected to sensorimotor regions of the brain and is more closely associated with normal motor functions.

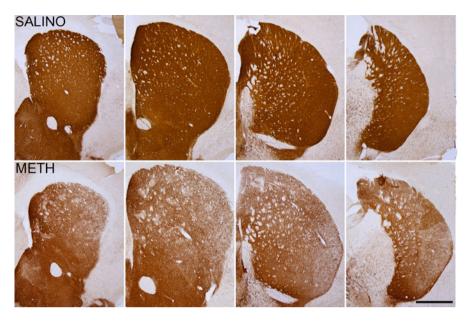
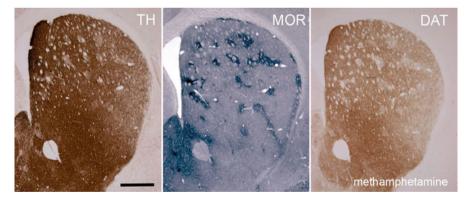
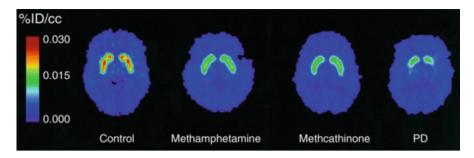


Fig. 3 Methamphetamine induces a loss of striatal TH-ir, indicative of dopamine fiber loss. Photomicrographs of striatal sections from mice treated with saline or methamphetamine stained for TH. METH produced a marked loss in striatal TH-ir that persists 7 days after treatment. Scale bar indicates  $500 \ \mu m$ 



**Fig. 4** Striatal vulnerability to methamphetamine. Methamphetamine toxicity occurs primarily in the striatum. Methamphetamine produces a preferential loss of TH and DAT in the striosomal compartment compared to the matrix. Photomicrographs of mouse brain sections 7 days after saline or methamphetamine treatment were stained for TH, MOR-1, and DAT. MOR-1 is a marker for striosomes. Scale bar indicates 500  $\mu$ m (Modified from Granado et al. (2010))

The pattern of dopamine degeneration in the striatum is similar to that observed in the early stages of other neurodegenerative diseases such as Huntington's disease, hypoxic ischemic injury, and treatment with MPTP, a selective neurotoxin for dopaminergic neurons that is frequently used as a model of Parkinson's disease.



**Fig. 5** Reduced DAT function in methamphetamine users. PET images showing accumulation of [11C] WIN-35,428 in the striatum of a control subject, an abstinent methamphetamine subject, an abstinent methcathinone subject, and a PD patient 70–90 min after injection of [11C] WIN-35,428 (Taken from McCann et al. (1998))

The nucleus accumbens is resistant to methamphetamine-induced dopaminergic axon loss (Granado et al. 2010), paralleling the effects of Parkinson's disease (Hurtig et al. 2000).

Methamphetamine neurotoxicity has also been demonstrated in humans (Fig. 5). PET studies in methamphetamine abusers found a reduction in DAT density in the caudate nucleus (26 % loss) and the putamen (21 % loss) after a short period of abstinence (Volkow et al. 2001a). Other authors reported similar DAT loss even 3 years after methamphetamine withdrawal (McCann et al. 1998). Studies in postmortem striatal tissue from chronic methamphetamine abusers showed a significant dopamine reduction concomitant with loss of DAT and TH immunore-activity, indicative of dopamine nerve fiber loss. These effects have been related to loss of neurological function, including memory loss and motor and verbal learning impairments (Volkow et al. 2001b). Methamphetamine also causes neurotoxicity in other brain areas including somatosensory parietal, frontal, and piriform cortex, olfactory bulb, and hippocampus, where apoptotic neurons have been found following exposure to the drug.

These studies in animals and in human abusers suggest that methamphetamine consumers may be more susceptible to neurodegenerative diseases like Parkinson's disease (PD), raising important concerns about the use and abuse of amphetamines clinically and recreationally. A very recent clinical study shows that methamphetamine users have a 76 % greater risk of developing PD than normal subjects. Although these studies have not yet confirmed neuronal damage in the substantia nigra of human methamphetamine abusers, results in animals support this loss and are consistent with the idea that repeated methamphetamine abuse predisposes the abuser to PD. Methamphetamine also causes neuronal loss in other brain areas such as the olfactory bulb, cortex, hippocampus, and striatum, as indicated by increased apoptosis, increased numbers of TUNEL-positive cells, and decreased numbers of neurons in the brains of methamphetamine-treated laboratory animals (Krasnova and Cadet 2009).

### 3 Mechanisms of Methamphetamine-Induced Neurotoxicity

#### 3.1 Role of Dopamine

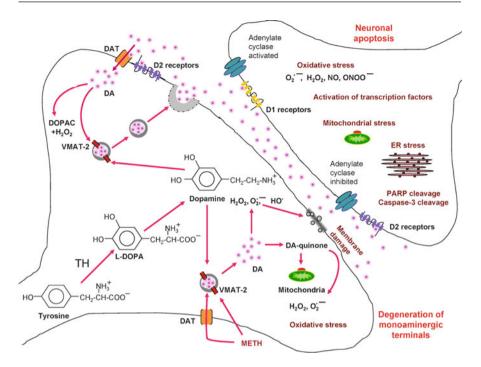
Methamphetamine consumption greatly increases the dopamine concentration in brain synapses. Excessive dopamine in the synaptic cleft is responsible for most of the physical and psychological effects of the drug, including addiction and psychomotor stimulant effects. An imbalance in the distribution of dopamine in the brain also seems to give rise to neurotoxicity, a fact that explains the localization of drug-induced degeneration to dopaminergic terminals (Fig. 6).

Following synthesis in dopaminergic neurons, dopamine is first released to the cytosol before being stored in vesicles where it is protected from metabolism and auto-oxidation. Methamphetamine induces a redistribution of dopamine inside the terminal, releasing dopamine from the vesicles into the cytosol where it is a substrate for metabolic and oxidative reactions, resulting in the production of dopamine quinones, superoxide anions, and hydrogen peroxide and hydroxyl radical species. This can further promote the oxidation of cytosolic dopamine, generating oxidative stress and leading to mitochondrial dysfunction and damage in the dopaminergic terminal (Cadet and Krasnova 2009; Thomas et al. 2008). The detrimental role of excessive cytosolic dopamine and its implication in the neurotoxic effects of methamphetamine is supported by the fact that when dopamine synthesis is inhibited by  $\alpha$ MPT, protection against methamphetamine toxicity is observed (Albers and Sonsalla 1995; Ares-Santos et al. 2012). Moreover, pretreatment with L-DOPA, a precursor of dopamine, and treatment with reserpine, which releases dopamine from vesicles to the cytoplasm, both potentiate methamphetamine toxicity (Albers and Sonsalla 1995; Granado et al. 2011a).

# 3.2 Implications of Oxidative Stress

Reactive nitrogen species and reactive oxygen species (ROS) are by-products of normal physiological metabolism in the brain, but excessive production of these reactive species can damage cell components, including lipids by lipid peroxidation, proteins by formation of protein carbonyls, and mitochondrial and nuclear DNA by peroxidation of these macromolecules. These reactive species impair mitochondrial respiratory chain enzymes and inhibit sodium-potassium ATPase, generating oxidative and nitrosative stress that leads to metabolic collapse and necrotic or apoptotic cell death. These oxidative stress cascades occur in several neurodegenerative disorders including Parkinson's disease.

Methamphetamine administration increases levels of extra-vesicular dopamine in the cytosol, where it can be metabolized by MAO or auto-oxidized in a process that generates toxic dopamine quinones. These dopamine quinones can damage cell proteins by binding to cysteine residues or by generating hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) and superoxide anions (O<sub>2</sub>•–), considered a major culprit in methamphetamine toxicity. The methamphetamine-induced increase in



**Fig. 6** Schematic representation of cellular and molecular events involved in methamphetamine-induced dopamine terminal degeneration and neuronal apoptosis within the striatum. The figure summarizes findings of various studies that have addressed the role of dopamine, oxidative stress, and other mechanisms in methamphetamine toxicity. Methamphetamine enters dopaminergic neurons via DAT and passive diffusion. Within these neurons, methamphetamine enters synaptic vesicles through VMAT2 and causes dopamine release into the cytoplasm via changes in pH balance. In the cytoplasm, dopamine auto-oxidizes to form toxic dopamine quinones, generating superoxide radicals and hydrogen peroxides via quinone cycling. Subsequent formation of hydroxyl radicals through interactions of superoxides and hydrogen peroxidative damage to presynaptic membranes. The toxic effects of released dopamine might occur through activation of dopamine receptors, as dopamine receptor antagonists block degeneration of dopamine terminals (Modified from Krasnova and Cadet (2009))

interaction of superoxides and hydrogen peroxide with transition metals like iron can lead to the formation of hydroxyl radicals (•OH) that cause oxidative stress, mitochondrial dysfunction, and peroxidative damage to dopaminergic terminal membranes (Krasnova and Cadet 2009).

The methamphetamine-induced increase in oxidative stress likely results from an imbalance between ROS production and the capacity of antioxidant enzyme systems to scavenge ROS: Methamphetamine both increases ROS production and reduces levels of the ROS scavengers CuZnSOD, catalase, glutathionine, and peroxiredoxins in the brain (Jayanthi et al. 1998; Li et al. 2008). Oxidative stress increases the susceptibility of the striosomal compartment to methamphetamineinduced dopaminergic toxicity (Granado et al. 2010). SOD is less abundant in striosomes than in the matrix (Medina et al. 1996), which may explain the greater vulnerability of the striosomes to methamphetamine neurotoxicity. This is supported by the finding that transgenic mice overexpressing CuZnSOD are resistant to methamphetamine-induced striatal neuronal damage (Hirata et al. 1996). Furthermore, antioxidants like ascorbic acid (vitamin C), vitamin E, bromocriptine (a hydroxyl radical scavenger), and coenzyme Q10 (antioxidant and mitochondrial energy enhancer) attenuate methamphetamine toxicity (Wagner et al. 1986). Free radical scavengers like PBN ( $\alpha$ -phenyl-*N*-terbutil nitrone) also reduce neurotoxic damage (Yamamoto and Zhu 1998), without altering the hyperthermic response that follows methamphetamine administration and that contributes to methamphetamine neurotoxicity (see Sect. 3.3 below).

# 3.3 Role of Hyperthermia

Methamphetamine usually produces a hyperthermic response in experimental animals and human meth abusers directly proportional to the dosage of the drug and to the ambient temperature in the place of treatment. This hyperthermia can be lethal (it is the first cause of methamphetamine-induced deaths) and can promote longterm neurotoxicity, as a relationship has been observed between the hyperthermic response and the neurotoxicity induced by the drug (Ares-Santos et al. 2012; Bowyer et al. 1994; Granado et al. 2010, 2011a). Strategies that reduce or avoid this hyperthermic response after methamphetamine treatment, like administering the drug at low ambient temperatures (4 °C) or pretreatment with pharmacologic agents like diclocipine or haloperidol, prevent or attenuate drug-induced neurotoxicity (Albers and Sonsalla 1995; O'Callaghan and Miller 1994). Furthermore, methamphetamine administration at high ambient temperatures promotes the hyperthermic response and increases neurotoxicity (Ares-Santos et al. 2012; Bowyer et al. 1994; Granado et al. 2011a; Miller and O'Callaghan 2003).

The correlation between hyperthermia and neurotoxicity is believed to be due to the fact that hyperthermia can potentiate DAT function (Xie et al. 2000), increasing free radicals and dopamine oxidation in the brain (Krasnova and Cadet 2009; LaVoie and Hastings 1999). Conversely, hypothermia has inhibitory effects on oxidative stress, as it reduces dopamine oxidation (LaVoie and Hastings 1999) and the formation of hydroxyl radicals (Fleckenstein et al. 1997; Krasnova and Cadet 2009). Moreover, neurons in animals maintained at low body temperature have reduced energetic demand, which may be protective since administration of meth-amphetamine produces striatal loss of ATP, possibly as a consequence of metabolic stress in dopaminergic neurons (Chan et al. 1994).

In contrast to these results, other pharmacological and genetic studies indicate that while hyperthermia contributes to methamphetamine-induced dopaminergic neurotoxicity, it is not required. For example, reserpine, a pharmacologic agent that dramatically lowers body temperature, strongly potentiates methamphetamine-induced neurotoxicity while blocking the hyperthermic response (Albers and Sonsalla 1995; Ares-Santos et al. 2012; Granado et al. 2011a). These results

indicate that blocking the hyperthermic response is not sufficient to protect against neuronal damage. Moreover, total or partial inactivation of DAT, nNOS, IL-6, or c-jun protects against methamphetamine-induced toxicity without altering the hyperthermic response. Thus, methamphetamine-induced hyperthermia contributes to, but is not required for, the neurotoxic effects of the drug.

#### 3.4 Role of Dopamine Receptors and Dopaminergic System

As evidence points to dysregulation of dopamine as the primary cause of methamphetamine-induced neurotoxicity, the role of the dopaminergic system in this process has also been evaluated. In particular, as methamphetamine acts as an indirect dopamine agonist, several studies have focused on elucidating the role of dopamine receptors. There are five different dopamine receptors (D1–D5), which fall into two families based on pharmacologic classification: the D1-like receptors (D1 and D5) and the D2-like receptors (D2, D3, and D4). Both families are involved in behavior and cognition, voluntary movement, motivation, punishment and reward, attention, working memory and learning, and in several neurodegenerative diseases like PD (Darmopil et al. 2009; Granado et al. 2008b; Ortiz et al. 2010).

Pharmacologic studies with the D2 receptor antagonists sulpiride, eticlopride, and raclopride have shown a dose-dependent prevention of methamphetamine toxicity in mice (Albers and Sonsalla 1995; Eisch and Marshall 1998). However, these compounds do not differentiate between members of the D2 receptor family, so it was not clear which receptor(s) mediate the protective effect. Recent studies using genetically modified mice lacking dopamine receptor D2 demonstrated that it is specifically the D2 receptor that is involved in methamphetamine toxicity as its genetic inactivation prevented the loss of dopaminergic striatal markers and inhibited the loss of dopaminergic neurons in the substantia nigra (Granado et al. 2011a).

D2 receptors are localized pre- and postsynaptically. At presynaptic locations, D2 receptors control dopamine release and thereby regulate extra-synaptic dopamine levels, which are involved in non-dopaminergic toxicity, such as in striatal medium spiny neurons, cortical and hippocampal neurons and neuropil. In addition, D2 receptors form heteromeric protein-protein complexes with DAT localized in the dopaminergic terminals that potentiate DAT activity. Blockade or inactivation of the D2R decreases striatal DAT activity. Since DAT knockout mice exhibit full protection against methamphetamine-induced dopaminergic toxicity (Fumagalli et al. 1998), indicating that active DAT is required for this neurotoxicity, the decrease in DAT is likely a major factor in the reduction of methamphetamine-induced dopaminergic toxicity induced by blockade or inactivation of the D2R. Moreover, fast scan cyclic voltammetry indicates that dopamine  $D2R^{-/-}$ mice have lower vesicular dopamine content, resulting in lower cytosolic dopamine levels. This also contributes to the reduction in methamphetamine-induced toxicity because cytosolic dopamine levels determine the severity of the toxicity (Granado et al. 2011a).

Receptors from the D1 family are also involved in methamphetamine-induced neurotoxicity as their pharmacologic inactivation with antagonists like SCH23390 also confers protection (Sonsalla et al. 1986). Genetic inactivation of the dopamine D1 receptor (D1R) also protected against reductions in striatal TH and DAT expression and against loss of dopaminergic neurons in the substantia nigra following methamphetamine administration, indicating that the D1R in particular is involved in methamphetamine neurotoxicity (Ares-Santos et al. 2012). Neuroprotection afforded by D1R inactivation is due in part to inhibition of hyperthermia, but also to the redistribution of dopamine inside the terminal. Animals lacking the D1R store more dopamine in vesicles and therefore have a reduced cytosolic dopamine pool compared to WT mice (Ares-Santos et al. 2012). Blockade of D1/D5R also suppresses activation of caspases 3 and 8, mediators of the calcineurin/NFAT/FasL-dependent apoptotic cell death pathway (Javanthi et al. 2005; Krasnova and Cadet 2009); this may also contribute to the neuroprotective effects of D1/D5 blockade or inactivation. Finally, SCH23390 decreases dopamine-induced oxidation and cytotoxicity mediated by ERK and JNK activation (Chen et al. 2004).

Other components of the dopaminergic system are also involved in methamphetamine-induced toxicity. Vesicular monoamine transporter (VMAT2) takes up dopamine from the cytosol to store it in synaptic vesicles, decreasing dopamine oxidation. Methamphetamine interacts with VMAT2 to cause a possible association of vesicles inside the dopaminergic terminal, increasing the release of dopamine to the cytosol and thereby increasing oxidative stress (Sulzer et al. 2005). VMAT2 knockout mice, with higher levels of cytosolic dopamine, are more sensitive to methamphetamine dopaminergic toxicity and show greater expression of oxidative stress markers than WT animals (Larsen et al. 2002). Other results are in line with these findings, showing that VMAT2 becomes nitrated 1 h after methamphetamine administration, which may reduce its activity (Eyerman and Yamamoto 2005), and that methamphetamine also reduces VMAT2 expression in the striatum (Krasnova and Cadet 2009).

### 3.5 Role of Glutamate and Nitric Oxide

Methamphetamine produces excitotoxicity by increasing glutamate release in the striatum (Nash and Yamamoto 1992), activating *N*-methyl-D-aspartate (NMDA) and alpha-amino-3-hydroxy-5-methyl-4-isoxazole propionic acid (AMPA) receptors. Stimulation of these receptors increases intracellular levels of Ca<sup>2+</sup>, causing activation of kinases, lipases, and proteases that damage the cytoskeleton, generating free radicals and DNA damage (Sattler and Tymianski 2000). Pharmacological studies using MK801, a noncompetitive NMDA receptor antagonist, prevented the long-term dopamine loss induced by methamphetamine (Sonsalla et al. 1989).

Moreover, NMDA receptor overactivation results in the production of superoxide radicals ( $O_2^{\bullet}-$ ) and nitric oxide (NO). When these two species react with each other, peroxynitrite (ONOO<sup>-</sup>), a more potent oxidative species, is formed, further potentiating neurotoxicity. To examine the contribution of this strong oxidizing agent to methamphetamine toxicity, a previous study measured the formation of neural 3-nitrotyrosine (3-NT), a product of tyrosine nitration that indicates irreversible structural modification of proteins (Butterfield et al. 2011) that can lead to loss of physiological cell functions, appearance of abnormal functions, and eventually cell death. A single injection of methamphetamine produced a significant rise in 3-NT concentrations in the striatum, signifying the involvement of ONOO<sup>-</sup> in the destructive effects methamphetamine abuse. Genetic or pharmacological inactivation of nNOS, the enzyme that produces nitric oxide in the brain, considerably reduced methamphetamine neurotoxicity without affecting the hyperthermic response (Itzhak et al. 1998, 2002), likely due to loss of peroxynitrite production.

### 3.6 Role of Astroglial and Microglial Activation

The central nervous system (CNS), consisting of the brain and the spinal cord, is an "immune privileged" area with an immune system distinct from that in the rest of the body. Microglia, a type of glial cell, are the resident macrophages of the central nervous system, and so act as its first and main form of active immune defense. These cells are normally in a resting state, but become activated after certain types of CNS damage, as a part of the innate immune response. Activated microglia migrate rapidly to the damage sites and secrete reactive species including proinflammatory cytokines (IL-1 $\beta$ , IL-6, TNF- $\alpha$ ) chemokines, prostaglandins, reactive oxygen species (ROS), nitric oxide, and superoxide to protect the brain. However, overactivation can be detrimental, producing cell death and astroglial dysfunction.

Amphetamines increase microglia activation in the striatum, hippocampus, cortex, and SN, with populations peaking 1 day after administration. Following administration of methamphetamine or other amphetamine derivatives, activated microglia are found in the areas in which dopaminergic neurotoxicity occurs, and the intensity of activation seems to be correlated with the level of dopaminergic damage. The highest levels of microglial activation occur in the dorsal striatum, an area highly affected by methamphetamine treatment, while the nucleus accumbens, more resistant than the striatum to methamphetamine-induced dopaminergic toxicity, has relatively few reactive microglia. Methamphetamine administration also results in increased levels of the three principal proinflammatory cytokines: IL-1 $\beta$ , IL-6, TNF- $\alpha$ , that in large part result from microglial activation (Clark et al. 2013). Thus, microglial activation represents a direct response to damage by amphetamines and is part of the cascade leading to neuronal damage (Thomas et al. 2008). Although many of the molecules secreted by activated microglia have been implicated in methamphetamine-induced neurotoxicity, and some anti-inflammatory drugs like ketoprofen, indometacin, tetracycline, and minocycline can protect against methamphetamine-induced microglial activation and neurotoxicity, attenuating microglial activation is not sufficient to protect against methamphetamine neurotoxicity (Sriram et al. 2006).

CNS damage is also accompanied by reactive gliosis, the injury-induced activation of astrocytes. The most abundant cells in the human brain, astrocytes, provide nutrients to nervous tissue, maintain extracellular ion balance, support other brain cells, and play an important role in repair and scar formation in the CNS after injury.

Recently, the immunomodulatory role of astrocytes has begun to emerge. Astrocytes can be protective, increasing levels of glutathionine (an antioxidant), facilitating sprouting, and providing growth factors, guidance molecules, and scaffolding for axonal regeneration, but they can also initiate several neuroinflammatory pathways and release inflammatory cytokines, some of which are neurotoxic. However, it seems likely that astrocytes play a positive role in limiting neuroinflammation and the balance between the activation of microglia and astrocytes leads to a detrimental or beneficial outcome (Clark et al. 2013).

Reactive gliosis is considered a universal reaction to CNS damage and is used as a sensitive marker of neuronal damage. Methamphetamine increases astrocyte activation in the striatum, as seen by increased expression of the marker glial fibrillary acidic protein (GFAP), which reaches maximal levels between 3 and 7 days after drug administration (Fig. 7). However, there are much earlier indications of reactive gliosis: Within a few minutes of methamphetamine delivery, there is already a 20 % increase in the magnitude of the Ca<sup>2+</sup> fluorescence signal in striatal astrocytes, and a 50–60 % increase in the number of responding astrocytes, indicating a primary astrocytic response (Granado et al. 2011b). As with microglia, activation of astrocytes takes place in the areas the most affected by methamphetamine, while the astrocyte population does not increase in the nucleus accumbens, where dopaminergic damage is normally not significant. Briefly, neuroinflammatory mechanism could in part contribute to the gradually escalating deleterious effects of methamphetamine.

# 3.7 Nrf2 and Inflammation Play a Role in Methamphetamine-Induced Neurotoxicity

The transcription factor Nrf2 (nuclear factor-erythroid 2-related factor 2) has been recently shown to have a protective role against methamphetamineinduced dopaminergic neurotoxicity (Granado et al. 2011b). Nrf2 is considered a master regulator of redox homeostasis, as it regulates the expression of a group of genes that encode the phase 2 detoxification enzymes, including heme oxygenase-1 (HO-1), NADPH quinone oxidoreductase (NQO1), and the catalytic and modulatory subunits of  $\gamma$ -glutamyl synthase (GCLM, GCLC) (Clark and Simon 2009; Johnson et al. 2008). In normal conditions, Nrf2 has a very short half-life because of its interaction with the BTB-Kelch protein Keap1, which promotes Nrf2 degradation by the proteasome (Lo et al. 2006). However, oxidant molecules are able to disrupt the Keap1/Nrf2 complex, rescuing Nrf2 from proteasomal degradation and allowing its entry into the nucleus and transcriptional activity.

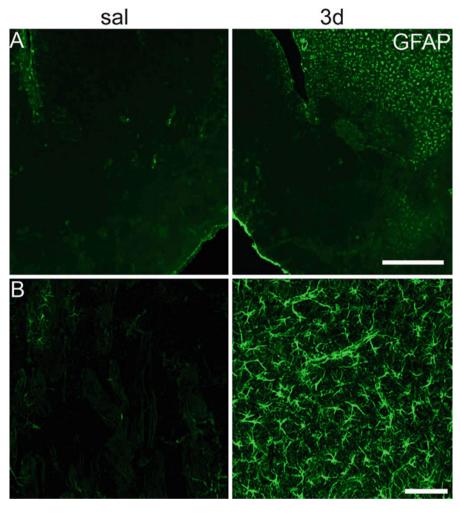


Fig. 7 Methamphetamine induces astrogliosis in mouse striatum but not in nucleus accumbens. (a) Photomicrographs of striatal sections from methamphetamine-treated mice stained for GFAP. (b) Enlargements of striatum sections shown in (a). Animals were killed 3 days after methamphetamine treatment. Methamphetamine increased GFAP staining in the striatum but not in nucleus accumbens. Bar indicates 500  $\mu$ m for (a) and 100  $\mu$ m for (b) (Modified from Granado et al. (2010))

Recent studies have shown that Nrf2 is activated by methamphetamine administration (Jayanthi et al. 2009) and that it plays a crucial role in the protection of dopaminergic neurons against oxidative stress by detoxifying mitochondrial complex I inhibitors and downregulating genes involved in the brain innate immune response. Mice lacking Nrf2 are more susceptible than WT mice to methamphetamine toxicity (Granado et al. 2011b), showing exacerbated hyperthermia, enhanced striatal TH and DAT-fiber loss, and greater decrease in dopamine levels, and increased dopamine and nigrostriatal dopaminergic alterations and gliosis following administration of methamphetamine (Granado et al. 2011b). This is due to the fact that in the absence of Nrf2, ARE-regulated genes in the striatum, including HO-1 and other antioxidant genes, are not induced by methamphetamine as they are in WT mice, leading to increased oxidative stress, accumulation of ROS (Chen et al. 2009), and ultimately to dopamine fiber loss. In addition, cytokine mRNA levels (TNF- $\alpha$ , IL-1 $\beta$ ), gliosis, and astrocytosis in the striatum were elevated to a greater extent in methamphetamine-treated Nrf2<sup>-/-</sup> mice than WT mice, indicating that the proinflammatory effects of methamphetamine treatment are potentiated in the absence of Nrf2.

Moreover, Nrf2<sup>-/-</sup> mice treated with methamphetamine presented significantly lower levels of modulatory subunits of  $\gamma$ -cysteine ligase (Gclm) and glutathione peroxidase (GPx) than WT animals, meaning that the detoxificating response is reduced in absence of Nrf2, a fact that could contribute to the increased detrimental effects induced by the drug.

Our finding that Nrf2-deficient mice were more sensitive than WT mice to methamphetamine-induced striatal damage further demonstrates that Nrf2 activation is part of a defensive response to methamphetamine neurotoxicity that involves modulation of methamphetamine-induced inflammation and oxidative stress (Granado et al. 2011b). This defensive modulation of inflammation and oxidative stress by Nrf2 is also seen following administration of other ROS-generating toxins like MPTP (Chen et al. 2009) and lipopolysaccharide (Rangasamy et al. 2004). Intriguingly, Nrf2 deficiency potentiates methamphetamine-induced neurotoxicity in the striatum but not in the SN. It is possible that methamphetamine differentially activates Nrf2-ARE transcription pathways in the striatum and the SN. Thus, differential regulation of Nrf2 by methamphetamine in SN and striatum might explain the lack of effect of Nrf2<sup>-/-</sup> on neurotoxicity in the SN in our study. These results strongly support the hypothesis that methamphetamine produces dopaminergic neurotoxicity through a process involving inflammation and oxidative stress (Granado et al. 2011b).

### 3.8 Role of Mitochondrial Dysfunction and DNA Damage

Methamphetamine-induced neurotoxicity also causes mitochondrial dysfunction and DNA damage. Mitochondria are the main source of cellular energy through activation of the ATP-producing mitochondrial respiratory chain, or electron transport chain, composed of a series of four complexes (I–IV). Methamphetamine is a cationic and lipophilic molecule that can diffuse into the mitochondria, where it is able to inhibit ATP synthesis (Asanuma et al. 2000; Krasnova and Cadet 2009). Administration of high doses of methamphetamine in rats decreases the activity of complexes II and IV of the respiratory chain in the striatum and prefrontal cortex, even in the absence of hyperthermia (Brown et al. 2005). Moreover, methamphetamine reduces ATP accumulation, resulting in mitochondrial dysfunction. The increase in reactive oxygen and nitrogen species may also contribute to the observed mitochondrial dysfunction.

Elevated production of oxygen- and nitrogen-based radicals and related non-radical products leads to the oxidation of essential macromolecules, including DNA. DNA damage plays a role in the pathogenic mechanism of methamphetamine, as the drug increases DNA oxidation in the striatum, hippocampus, substantia nigra, and olfactory bulb (Jeng et al. 2006), causing apoptotic cell death in experimental animals (Deng and Cadet 2000). In particular, the 8-oxoguanine (8-oxoG) produced in DNA lesion may mispair with adenine, causing transversions or mutations, altering the DNA binding of nuclear transcription factors or blocking RNA polymerase, resulting in altered or delayed transcription of proteins. DNA repair genes, including members of the BER pathway, are upregulated in adult mice after methamphetamine administration, suggesting that increased repair activity is induced to counteract the oxidative DNA damage induced by the drug.

# 4 Neuroprotective Strategies Against Methamphetamine-Induced Neurotoxicity

As basic and clinical research findings begin to elucidate mechanisms of methamphetamine-induced neurotoxicity, potential neuroprotective strategies are being proposed. Since oxidative stress and mitochondrial dysfunction are important factors in methamphetamine-induced neurodegeneration, pretreatment with antioxidants like *N*-acetyl-L-cysteine, ascorbic acid, vitamin E, or coenzyme  $Q_{10}$  was evaluated and shown to have protective effects against depletion of monoaminergic axons (Krasnova and Cadet 2009). Melatonin, another antioxidant compound that also has antiapoptotic effects, also reduces methamphetamine-induced depletion of dopaminergic markers in the striatum (Hirata et al. 1998).

Other strategies for avoiding an increase in oxidative stress, such as preventing methamphetamine-induced hyperthermia by pharmacologic treatment or by maintaining animals at cool ambient temperatures during drug administration, also reduce mortality and neurotoxicity.

Controlling the amount of dopamine in the cytosol, where it is susceptible to oxidation, causing oxidative stress, is another neuroprotective strategy. Increasing VMAT2, the vesicular monoamine transporter that sequesters dopamine into vesicles, and administering dopamine uptake inhibitors or dopamine receptor antagonist, all provide protection against methamphetamine -induced degeneration of striatal dopamine terminals. In addition, some trophic factors provide protection against the toxic effects of methamphetamine. Administration of glial cell line-derived neurotrophic factor (GDNF) or brain-derived neurotrophic factor (BDNF) can prevent methamphetamine-mediated reduction in dopamine in the striatum and caspase activation (Dluzen 2004; Matsuzaki et al. 2004; Melega et al. 2000).

Since neuroinflammation may contribute to methamphetamine neurotoxicity, some cytokines, such as interferon gamma, TNF-alpha among others, have been

evaluated and were shown to protect against methamphetamine toxicity. Estrogen can also protect against methamphetamine-induced damage to the nigrostriatal dopamine system (D'Astous et al. 2005). Pretreatment with tamoxifen, a selective estrogen receptor modulator, showed neuroprotective effect against methamphetamine-induced toxicity and attenuated inflammatory response in the striatum when used alone but abolishes estrogen's positive effects when combined with this hormone. While both treatments prevented dopamine decrease, estrogen protected more efficiently other dopaminergic parameters, suggesting that overall estrogen is more effective than tamoxifen as a neuroprotectant of the nigrostriatal dopaminergic system (Bourque et al. 2007; D'Astous et al. 2005).

#### 5 Conclusion

Methamphetamine is a synthetic drug used worldwide, mostly for recreational purposes, due to its powerful psychostimulant effects. It has addictive effects due to its structural analogy with dopamine, but also has neurotoxic effects on the dopaminergic system. The drug causes a reduction in dopamine markers in the striatum similar to that seen in the early stages of neurodegenerative diseases like PD, Huntington's disease, and hypoxic/ischemic injury. In experimental animals, methamphetamine also causes neuron loss in the substantia nigra. This may explain why patients who abuse methamphetamine have a predisposition to future development of PD. Among the mechanisms responsible for methamphetamine's neurotoxic effects are oxidative stress, hyperthermia, microglia and astroglia activation, mitochondrial dysfunction, DNA damage, and elevated levels of glutamate and nitric oxide. The recent implication of the D1 and D2 receptors in methamphetamine-induced neurotoxicity suggests that targeting these receptors may be a promising strategy for development of new approaches to prevention and treatment of methamphetamine addiction and its neurotoxic effects. In addition, the neuropathological similarities between methamphetamine neurotoxicity and PD and the demonstrated predisposition of methamphetamine abusers to developing PD indicate that similar therapeutic approaches may be useful in the early stages of PD and related neurodegenerative diseases.

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# Pathophysiology of Obsessive-Compulsive Disorder: Insights from Normal Function and Neurotoxic Effects of Drugs, Infection, and Brain Injury

Henry Szechtman, Sohel Shivji, and Erik Z. Woody

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# Abstract

The clinical presentation of obsessive–compulsive disorder (OCD) is summarized and a theoretical model of the disorder and its neurobiology are described. The model is used as a framework to account for the observed pattern of clinical symptoms, for the pathophysiology of OCD, and for medical reports of OCDlike symptoms from neurotoxic effects of drugs, brain injury, and infection. The essential clinical feature of OCD is preoccupation with ideas and behaviors called "obsessions" and "compulsions," respectively. The characteristic content of obsessions and compulsions is concern about dangers that might happen in the

H. Szechtman (🖂) • S. Shivji

Department of Psychiatry and Behavioural Neurosciences, McMaster University, Hamilton, ON, Canada

e-mail: szechtma@mcmaster.ca

E.Z. Woody Department of Psychology, University of Waterloo, Waterloo, ON, Canada e-mail: ewoody@uwaterloo.ca

future – as opposed to actual threats that are direct and immediate. Hence, OCD is a clinical preoccupation with *potential* threats. A model is considered where normal concerns regarding potential danger and security are handled by a biologically ancient hard-wired motivational system, called security motivation system (SMS), which has as its output the same repertoire of precautionary behaviors that characterize OCD compulsions. In the model, OCD symptoms emerge because security motivation is normally deactivated by a negative feedback signal generated by performance of precautionary behaviors, but in patients with OCD, this signal is inoperative, resulting in continued motivation to perform security-related behaviors. The neurobiology of SMS consists of functional loops involving a cascade of cortico-striato-pallido-thalamo-cortical connections, with inhibitory connections from the brainstem. Accordingly, OCD pathophysiology is overly persistent and uncontrolled neural activity in SMS, possibly due to a dopamine-serotonin imbalance. Evidence of OCD symptoms from drugs, brain injury, and infection is consistent with disturbed basal ganglia regulation as the pathophysiology of OCD.

# 1 Introduction

A comprehensive understanding of the pathophysiology of obsessive–compulsive disorder (OCD) is lacking because the cause of this disease is unknown. In general, if the cause of a disease is known then the course of pathogenesis can be traced forward from a well-defined starting point. In this way, even when the pathophysiology and presentation of symptoms are varied, it is nevertheless certain that they all belong to the same disease (stem from the same cause). However, if the disease cause is unknown, and one observes variations in pathophysiology and symptoms, it is not certain whether those variations belong to one disease with several presentations or in fact reflect several different diseases with different causes.

Criteria for disease causality were laid down by Robert Koch in 1882 when he proved the causation of tuberculosis, a devastating disease then believed to result from societal woes (Barnes 2000). Koch established that a bacterium (which he called "tubercle bacillus") is the causal agent of tuberculosis and he proved this through experiments demonstrating:

(1) The parasite occurs in every case of the disease. (2) The parasite does not occur in other diseases or nonpathogenically. (3) After being fully isolated and repeatedly grown in pure culture the parasite can induce the disease by being introduced into a healthy animal. (Barnes 2000, p. 433)

As noted, when it comes to OCD, its cause – its agent of origin – is unknown. In fact, whether there are multiple sufficient causes that result in OCD is also unknown. However, information exists regarding the nature of the behavioral systems affected by OCD, and this information can be integrated into a theoretical model that provides a useful framework for considering the pathophysiology of OCD and how the function of these systems can be affected by neurotoxicity.

Below, the clinical presentation of OCD is summarized and a theoretical model is described that provides a framework to explain the symptoms of OCD and their neurobiological underpinnings. This framework is followed by a review of case reports to illustrate the neurotoxic effects of drugs, brain injury, and infection as causal agents producing OCD-like behavior.

# 2 Description of OCD

Given that the cause and pathophysiology of OCD are unknown, there are no biological tests for OCD, and, hence, a diagnosis of OCD is made solely from evidence of a particular behavioral profile (American Psychiatric Association. Task Force on DSM-IV 2000). The essential features of this behavioral profile are preoccupation by the individual with certain types of ideas and behaviors. The preoccupation is clinically significant because it is (a) excessive (it consumes more than 1 h in a day and interferes with normal daily function), (b) ego-dystonic (it gives no pleasure to the individual but the subject feels driven to be so engaged), and (c) unreasonable (the person acknowledges the preoccupation may appear to others as disconnected from objective reality). The ideas and behaviors that preoccupy the individual are called "obsessions" and "compulsions," respectively. According to Reed (Reed 1985), the content of obsessions may include thoughts involving (a) self-depreciation, self-denigration, and unworthiness either with reference to perceived social norms or the individual's own standards; (b) undue concern with one's bodily functions; (c) fears, doubts, and preoccupations about dirt and the spread of disease to self; and (d) fears of harm to a friend or relative and being responsible for the imagined event. The range of compulsive behaviors includes (Reed 1985) (a) excessive *checking* activities, characterized by repeated re-doings of actions supposedly related to security, orderliness, or accuracy; (b) avoidance behaviors, which are "activities engaged in to avoid feared objects, places, or situations" (p. 37); and (c) washing and cleaning, generally of hands but sometimes also compulsive washing of clothes, teeth cleaning, or the cleaning of possessions or parts of the home. Individuals may be preoccupied with more than one obsession and/or compulsion. The two most common characteristics of obsessions are doubts and indecision, and the two compulsive behaviors performed most often are checking and washing (Rasmussen and Eisen 1992).

It is remarkable that despite being aware that their preoccupation is exaggerated, afflicted individuals feel unable for any extended period of time to control their obsessions and compulsions; indeed, powerlessness over time spent engaged in obsessions and compulsions is very often the immediate reason for seeking help. But OCD does not reflect weaknesses of the will (Huoranszki 2011) or intellect. Most revealingly, there are individuals with OCD who despite the distresses of their illness manage very prolific and intellectually exceptional careers, one famous example being Samuel Johnson, the eighteenth century English writer and literary critic who counts among his major works the sole authorship of "A Dictionary of the English Language." There are many such examples of highly successful individuals with OCD where clearly neither weak will nor poor intellect are the reasons for OCD symptoms (the interested reader may view video testimonials and descriptions of OCD on YouTube, e.g., http://www.youtube.com/watch?v=Rn1OYIYzgm8 or http://www.youtube.com/watch?v=dSZNnz9SM4g). Rather, the striking disconnection in OCD between knowing rationally that some actions are unreasonable, but nevertheless feeling that they must be performed, demonstrates the presence of a powerful "feeling" system as a determinant of behavior (Kahneman 2011), a notion which is elaborated upon in the next section.

It is should be noted that the DSM-IV manual indicates criteria for the differential diagnosis of OCD. Specifically, it states that OCD "must be distinguished from Anxiety Disorder Due to a General Condition" or from a "Substance-Induced Anxiety Disorder." Those diagnoses apply, respectively, "when the obsessions or compulsions are judged to be a direct physiological consequence of a specific general medical condition" or "when a substance (i.e., a drug of abuse, a medication, or exposure to a toxin) is judged to be etiologically related to the obsessions or compulsions" (American Psychiatric Association. Task Force on DSM-IV 2000). At first glance, the differential diagnosis suggests the presence of another disease, with a pathophysiology distinctly different from the pathophysiology of OCD diagnosed solely from behavioral evidence. However, the utility of the differential diagnosis probably applies to the clinical management of the symptoms rather than indicating that the differently labeled disorders have their own distinct pathophysiology. Indeed, as argued later, the medical conditions producing compulsions and obsessions reveal a plausible pathophysiology of OCD.

# 3 Security Motivation System as the Framework to Explain OCD

Disease in an organism was seen for a long time as due to invasion by unseen spirits, and it is relatively recently that the unseen forces have been given a physical reality, as demonstrated by Robert Koch above. The history of psychiatric disorders is no different, and even today mental illness is often viewed as a creation of outside forces, such as possession by various spirits or demons, environmental evils, abnormal learning, or mutation of genes. Nevertheless, it is crucial to appreciate that whatever the actual agent may turn out to be, and regardless of how bizarre the psychiatric symptoms may seem, the psychiatric condition is not a de novo creation but a phenomenon already in the repertoire of the organism. After all, the agent must act on the body and consequently the observed behavioral output, regardless of how bizarre, is a reflection of the functions the brain is built to perform. Accordingly, psychiatric disorders, including OCD, reflect the exaggerated operation of normal brain system(s) present in the population. This perspective raises the following two defining and related questions regarding OCD:

(i) Is there a common theme to the symptoms of OCD, or are the symptoms merely a collection of unrelated acts?

(ii) What are the normal constitutive processes that can lead an individual to feel driven to engage in irrational acts (obsessions and compulsion) but without showing a disconnection from objective reality?

With regard to a possible relationship among OCD symptoms, a perusal of the various obsessions and compulsions (Reed 1985) suggests that they can be conceptualized as belonging to a specific domain. In particular, the content of most OCD thoughts and behaviors seems to revolve around concerns related to security or safety, either of the self or of others. Hence, following an expression used by Paul MacLean (MacLean 1973), the core theme of OCD obsessions and compulsions may be described as concerns regarding "self-preservation or preservation of the species" (Szechtman and Woody 2004). Importantly, because the concerns in obsessions and compulsions are about dangers that *might* happen in the future – as opposed to actual threats that are direct and immediate – it is *potential* threats that are proposed to constitute the domain of OCD concerns (Szechtman and Woody 2004; Woody and Szechtman 2011).

The domain of potential threats belongs to the normal concerns of everyone and is not unique to OCD (Muris et al. 1997; Rachman and De Silva 1978; Rassin and Muris 2007; Reed 1985; Salkovskis and Harrison 1984). Within this normal domain several classes of potential threats are present, but of most salience to OCD are the security concerns engendered by predation (attack by others) and contamination–contagion. Interestingly, these security concerns are among the key normal fears of children, with a clear developmental time-course (Boyer and Bergstrom 2011). The two classes of potential dangers normally evoke appropriate and specific precautionary responses, namely, checking and washing, respectively. As noted, the exaggerated performance of checking and washing are the two most common OCD compulsions (Rasmussen and Eisen 1992). Conceivably, OCD subtypes of checkers and washers map onto the predation and the contamination–contagion classes of security concerns, respectively.

There are crucial distinctions between threats that are potential and those that are direct and immediate, with profound implications for the design of systems that process them (Boyer and Lienard 2006; Szechtman and Woody 2004; Woody and Boyer 2011; Woody and Szechtman 2011). One important distinction is that the two types of threat necessitate different types of responses. Already-present threats require fast responses to confront the danger, the faster the better – whereas potential ones require behavioral adjustments that can take more time to develop and perform. Furthermore, behaviors evoked by immediate threats are directed and constrained by moment-to-moment interaction with the source of danger. In contrast, potential threats are countered by indirect precautionary measures, such as avoidance or probing, and are free of constraint by an actual confrontation.

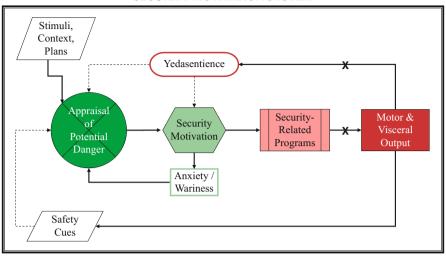
Another important distinction between potential and immediate threats relates to feedback from external environment regarding elimination of the threat. With actual threats, cues of danger confront the senses and thus there is no ambiguity about the presence of the threat or its successful elimination. In contrast, for potential threats there is an information asymmetry regarding presence versus elimination of threat. Potential dangers such as contagion or predator presence are detected on the basis of particular partial cues, such as rotting smells or predator spoor. However, once the organism has engaged in precautionary behaviors, there is no positive evidence that these did actually eliminate or sufficiently reduce the danger. For example, the absence of a predator smell is no positive evidence for the predator's absence. Thus, taking precautions does not create clear signals that the danger is removed. Hence, with potential threats, the external environment provides no actual feedback either that the danger is eliminated or that the precautionary behavior was appropriate and effective.

This peculiar property inherent to potential threat – no feedback from external information that performance of precautionary behaviors achieved security – creates an obvious design challenge for a system handling potential threats: (1) what ensures that precautionary behaviors are initiated and performed and (2) what decides that a satisfactory level of security has been achieved? As explained below, it is hypothesized that this challenge is met through a biologically "hard-wired" special motivation – the security motivation system (SMS) (Szechtman and Woody 2004, 2006; Woody and Szechtman 2005, 2011).

The challenge of ensuring that precautionary behaviors are initiated and performed is answered by employing a dedicated motivational system. A defining design property of motivation is the urge to engage in species-typical preprogrammed responses when the motivation is activated. Accordingly, a mechanism to process potential threats through a dedicated security motivation system has a built-in urge to perform security-related precautionary behaviors in response to potential threats.

The next challenge – how to satisfy security without external feedback – is answered by utilizing signals generated internally. Most motivations are deactivated through contact with the appropriate goal object; for example, hunger motivation is deactivated by eating food, which generates a negative feedback satiety signal that terminates the motivation to attain food (Toates 1986). Because no external goal objects exist to satisfy an activated security motivation, then it must be that motivation gets deactivated through the very act of performance of security-related precautionary behaviors. In other words, self-generated signals from engagement in precautionary behaviors serve as the negative feedback satiety signal that terminates security motivation. Such an arrangement is functionally plausible because species-typical motor activity per se is rewarding (Glickman and Schiff 1967). Clearly, the built-in reward for security-related engagement contributes also to maintenance of precautionary responses to potential threats.

The model of SMS shown in Fig. 1 makes comprehensible how OCD can represent the exaggerated activity of a normal system and how normal constitutive processes can indeed compel performance of seemingly irrational acts (obsessions and compulsion) despite awareness of a seemingly inconsistent objective reality. The design in Fig. 1 of SMS follows the generic blueprint for a special motivation and consists of four modules shown in the middle row: a specialized module for detection of potential threats (*green* circle labeled "appraisal of potential danger") that turns on a "security motivation" module, which generates a coordinated set of processes to energize and focus brain resources on attaining a specified goal.



**Fig. 1** A conceptual model of the security motivation system and sites of dysfunction producing OCD. Solid arrows indicate excitatory and dashed arrows inhibitory stimulation, respectively. Yedasentience output does not act on environmental input but rather on the appraisal of potential danger and the security motivation processors to inhibit their activity. Exposure through motor output to "safety" stimuli provides inhibitory stimulation to appraisal of potential danger [Mod-ified from (Szechtman and Woody 2004)]

The signal from "security motivation" selects from the "security-related programs" module the plan of programs appropriate for the particular class of potential threat (e.g., washing and cleaning behaviors for the contamination–contagion class, probing and checking for the predation class). The "security-related programs" module in turn activates the "motor and visceral output" module, which executes and implements the actual performance of the species-typical motor actions. Execution of precautionary responses generates also internal feedback signals, including the signal labeled "yedasentience" shown in red in the top row [*yedasentience* = "feeling of knowing" (Szechtman et al. 2002; Szechtman and Woody 2004; Woody and Szechtman 2000)]. This yedasentience signal inhibits the "security motivation" and "appraisal of potential danger" modules, thereby stopping SMS activity.

Termination of security motivation activity by the self-generated yedasentience negative feedback signal provides an explanatory mechanism for OCD compulsions and obsessions. Specifically, failure to generate yedasentience (or respond to this signal) leaves SMS activity on for a prolonged period of time, driving persistent performance of precautionary behaviors – this continued performance constitutes OCD obsessions and compulsions. Two potential sites where damage may lead to a failure to generate yedasentience and yield obsessions and compulsions are shown by X in Fig. 1.

#### SECURITY MOTIVATION SYSTEM

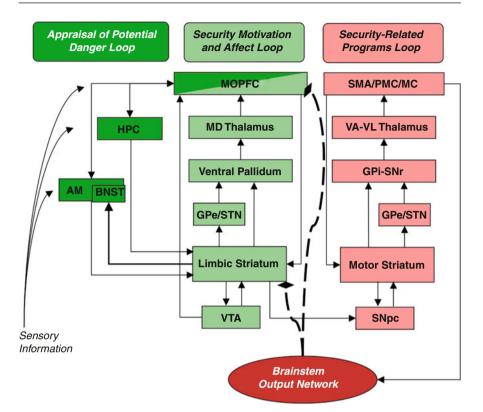
Lack of yedasentience feedback also explains how seemingly unreasonable obsessions and compulsions can exist alongside a normal sense of reality. Two particular characteristics of SMS contribute to the explanation. The first relevant property is the hierarchical dominance of an activated SMS. Specifically, when SMS is activated, it functions at the top of the hierarchy among competing systems controlling brain resources and actions. This is a design feature of special motivations which are engaged to perform functions vital for survival of the organism (Hebb 1966). Hence, it follows that when security motivation is stimulated, the individual becomes preoccupied with security, at the expense of other cues in the environment, and in this respect such preoccupation is normal and does not reflect any loss of a sense of reality.

The second relevant property is peculiar to SMS: an inherent disconnection of SMS deactivation from objective reality. As noted above, the external environment provides no reliable feedback either that potential threat was eliminated or that precautionary behaviors were appropriate and effective in achieving security. Consequently, as shown in the model in Fig. 1, information from the external environment has no access to security motivation to shut it down – only the self-generated yedasentience feedback can turn off security motivation. Hence, the drive to engage in obsessions and compulsions can coexist with a logically conflicting sense of reality.

# 4 Neurobiology of SMS and Substrate of OCD Pathophysiology

Framing OCD as a disturbance of security motivation simplifies the task of identifying OCD pathophysiology because it reduces the pathophysiology of OCD to a dysfunction in the neurobiology of SMS, and a fairly good model of a functional SMS neurobiology exists (Szechtman and Woody 2004, 2006; Woody and Szechtman 2011). The model follows the generic plan of motivational systems derived from behavioral neuroscience studies on motivation. Accordingly, the proposed schematics of SMS neurocircuitry and physiology are shown in Figs. 2 and 3.

Like other neuroanatomical models of motivation [e.g., (Brown and Pluck 2000; Everitt and Wolf 2002)], the SMS diagram in Fig. 2 consists of functional loops involving cortico-striato-pallido-thalamo-cortical connections, as advanced by Alexander et al. (1986) and others (Groenewegen et al. 1999; Haber et al. 2000; Joel and Weiner 2000; Penney and Young 1983). Importance of the yedasentience signal is highlighted by inhibitory connections from the brainstem. There are four cascading neural loops in Fig. 2, each mediating one of the four functional components in Fig. 1 – the color codes of the loops correspond to the functional modules in Fig. 1. Accordingly, the circuits in Fig. 2 are labeled: appraisal of potential danger loop (*green*), security motivation and affect loop (*light green*), securityrelated programs loop (*light red*), and brainstem output network (*red*), where each loop contains specific brain structures that are interconnected to form a specialized neurocircuit mediating a distinct motivational task.



**Fig. 2** A neural circuit model of the security motivation system. Each of the four distinct subcircuits (*loops*) subserves one of the functional components in Fig. 1 and is identified by corresponding colors. The dashed line indicates possible sites of yedasentience feedback inhibition. Abbreviations: *AM* amygdala, *BNST* bed nucleus of the stria terminalis, *GPe* external segment of the globus pallidus, *GPi* internal segment of the globus pallidus, *HPC* hippocampus, *MC* motor cortex, *MD* Thalamus, mediodorsal thalamic nucleus, *MOPFC* medial prefrontal cortex and orbital prefrontal cortex, *PMC* premotor cortex, SMA, supplementary motor area, *SNpc* substantia nigra pars compacta, *SNr* substantia nigra pars reticulata, *STN* subthalamic nucleus, *VA* ventroanterior thalamic nucleus, *VL* ventrolateral thalamic nucleus, *VTA* ventral tegmental area [Reprinted by permission from (Szechtman and Woody 2004)]

The first motivational task – detection of potential danger – is handled by the appraisal of potential danger loop composed of the interconnected green-coded brain regions in Fig. 2. The indicated interconnected limbic regions – the hippocampus, the amygdala, and the bed nucleus of the stria terminalis (BNST), as well as the medial and orbital prefrontal cortex (MOPFC) – process motivation-related input, including security-related information (Bechara et al. 2000; Fiddick 2011; LeDoux 2000; Woody and Szechtman 2011). Thus, this detection circuit has the computational functions to quickly process signalling cues (e.g., rotting smells, predator spoor, novelty) and more complex information ("context," "plans") and determine whether a change of present circumstance could result in a future harm. If computational outcome is indeed potential danger, the appraisal of potential danger loop has neuroanatomical links needed for activation of the next motivational task, handled by the security motivation and affect loop (*light green*-colored boxes). Specifically, the hippocampus and the amygdala have projections to the ventral striatum, a key area in the security motivation and affect loop; moreover, the MOPFC (denoted by the box with *green* and *light green* colors) is a component of both the appraisal of potential danger loop and the security motivation and affect loop, providing an interface between the two circuits.

The next motivational task is the engagement of motivational drive and affect and is handled by the security motivation and affect loop (light green-coded brain regions in Fig. 2). The neuroanatomical sites of this loop are linked as a prototypical basal ganglia-thalamocortical circuit (Alexander et al. 1986) and form, specifically, the "limbic striatum loop" (Brown and Pluck 2000; Joel and Weiner 2000). Three attributes of this neurocircuit seem especially relevant for the designated job of the security motivation and affect loop. First, it contains dopamine innervation from the VTA onto the limbic striatum and the prefrontal cortex. Dopamine input can supply the tonic ignition of the loop and this dopamine stimulation is likely the implementation of a salient characteristic of motivations, namely, the urge to engage in goaldirected behaviors. Indeed, stimulation of the mesolimbic dopamine system yields approach and investigatory behaviors (Ikemoto and Panksepp 1999), and such motor activity is part of the repertoire of many motivations, including security motivation, where it manifests itself as probing the environment for potential danger. Second, the loop involves connections with limbic regions implicated with processing of affect accompanying motivation. In the case of security motivation, it has been suggested that the connection from the limbic striatum to BNST (Davis and Shi 1999; Davis et al. 1997; Lang et al. 2000; Woody and Szechtman 2011) is especially important in mediating the feelings of anxiety and wariness, the phenomenological cues of potential danger (Masterson and Crawford 1982; Woody and Szechtman 2011). Finally, the neural arrangement of the regions within the loop is such as to prolong and sustain reverberatory activity (Alexander et al. 1986; Penney and Young 1983) needed to maintain motivational drive and also activate the next motivational task.

The next motivational task is the selection and engagement of an appropriate security-related response. This job is executed by the security-related programs loop (light-red-coded brain regions in Fig. 2); this loop is also a prototypical basal ganglia neurocircuit but interconnected with the dorsal (or "motor" striatum) and innervated by dopamine from the substantia nigra. The circuit has design features especially useful for "selection" of appropriate responses (Mink 1996; Penney and Young 1983; Wolgin 2012). Every motivation has its own set of species-typical behavioral actions; typically, the neural instructions of how to implement the organization of a species-typical pattern of behavior are held in the striatum in the form of a precoded "program" (Aldridge et al. 2004; MacLean 1978; Penney and Young 1983; Wise and Rapoport 1989). Accordingly, depending on the domain of potential danger, the security-related programs loop engages the appropriate program (e.g., for probing or checking the environment in case of potential threat of

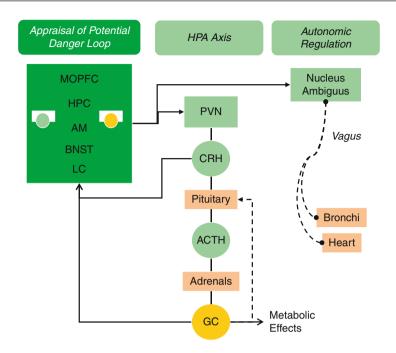
attack and for washing or grooming in case of contamination–contagion), and at the same time it inhibits and prevents competing behavioral programs from access to motor output. Dopamine activity from the nigrostriatal system probably helps to maintain the neural operation of the selected program and intensity of its execution, reflected in the vigor of motor performance (Dvorkin et al. 2010).

Finally, the program selected by the security-related programs loop implements those instructions by activating the brainstem output network (*red*-coded area in Fig. 2), and stimulating brainstem output neurons that convey neural signals for the actual contraction of muscles that move the body and viscera in performance of the species-typical actions. In addition to patterned movements, the brainstem output network generates the yedasentience negative feedback signal, which shuts down security motivation via inhibitory input possibly to the ventral striatum and MOPFC (heavy dashed line in Fig. 2). This negative feedback pathway is probably serotonergic, given that the serotonin neurotransmitter serves widely as a "stop" or "satiety" signal in many functions (e.g., Blundell 1991; Lorrain et al. 1999; Soubrie 1986) and serotonin reuptake inhibitors have therapeutic effects in OCD (DeVeaugh-Geiss 1994).

Every motivation which has a characteristic neural network possesses also a characteristic physiology that facilitates network operation at its optimum. The model in Fig. 3 shows that physiological mechanisms that support an activated security motivation operate through stimulation of the hypothalamic–pituitary– adrenocortical (HPA) axis and regulation of the parasympathetic nervous system (PNS). These systems contribute peripherally and centrally to enhance security motivation, which is turned on by potential threat and thus geared for action – for doing *something* that will relax the psychological tension of uncertain danger. To support those SMS prerequisites, HPA and PNS engage peripheral mechanisms to mobilize energy for physical work and also exert central effects on the *appraisal of potential danger* and *security motivation and affect* loops to potentiate threat detection and motivation [Fig. 3; for details, see (Woody and Szechtman 2011)].

### 5 Pathophysiology of OCD

Using the above framework of a fully functional SMS (Figs. 2 and 3), the pathophysiology of OCD is relatively clear cut – OCD pathophysiology is normal neural activity in SMS loops that becomes overly persistent and uncontrolled (Saxena et al. 1998; Wise and Rapoport 1989). While there are many possible neuroanatomical spots where a malfunction (regardless of cause) can produce this pathophysiology, two salient malfunctions likely to yield prolonged and uncontrolled SMS neural activity are excessive dopaminergic stimulation, insufficient serotonergic negative feedback, or possibly both. Indeed, there is evidence for each possibility in OCD (Denys et al. 2004; Westenberg et al. 2007), including dysfunction in both dopamine and serotonin (Nikolaus et al. 2010) attested by current clinical practice of combining dopamine blockers with serotonin reuptake inhibitors for treatment of OCD (Bloch et al. 2006; Dougherty et al. 2004).



**Fig. 3** The physiological network associated with security motivation. The appraisal of potential danger loop of Fig. 2 connects with the HPA Axis, which has both input from and output to the Appraisal Loop. Within the Appraisal Loop there are receptors for CRH and GC, denoted by the respective circles in white rectangles, by which HPA can modulate it; CHR also acts as a neurotransmitter within the Appraisal Loop. Finally, the Appraisal Loop affects Autonomic Regulation, by which the myelinated vagus modulates the bronchi and heart. Abbreviations: *AM* amygdala, *BNST*, bed nucleus of the stria terminalis, *CRH* corticotropin-releasing hormone, *GC* glucocorticoids, *HPC* hippocampus, *LC* locus coeruleus, *MOPFC* medial prefrontal cortex and orbital prefrontal cortex, *PVN* paraventricular nucleus [Reprinted by permission from (Woody and Szechtman 2011)]

One manner by which dopamine–serotonin imbalance can yield excessive SMS activity and thus OCD pathophysiology is as follows. OCD concerns are the normal security-related concerns of everyone and indeed their incidence does not differ between OCD and non-patient populations (Muris et al. 1997; Rachman and De Silva 1978; Rassin and Muris 2007; Reed 1985; Salkovskis and Harrison 1984). Consequently, individuals are often presented with some potential threat that activates the dopaminergic motivational drive of the security motivation circuit. However, in individuals with a constitutively hyperfunctional dopamine system, activation of security-related dopaminergic motivational drive produces excessive stimulation of dopamine receptors thereby driving the SMS circuit continually and overpowering any negative feedback to shutdown SMS activity. Similarly, in individuals with a constitutively ineffective serotonergic negative feedback,

activation of security motivation also yields OCD pathophysiology but here SMS activity stays prolonged not because of an intrinsic or pathologic dopamine hyperfunction. Instead, SMS activity is prolonged because serotonergic negative feedback is the normal mechanism to deactivate the dopaminergic motivational drive, and consequently, without a serotonergic "stop" signal, SMS activity continues unabated.

# 6 Evidence for Overactive SMS Circuit in OCD

The foundational line of evidence for overactive functional loops involving corticostriato-pallido-thalamo-cortical connections came from early positron emission tomography (PET) studies measuring the brain distribution of glucose metabolism in patients with OCD. Such PET studies showed metabolic hyperactivity in orbitofrontal cortex and caudate nucleus in patients with OCD, a hyperactivity that resolved upon disappearance of OCD symptoms with therapy (Baxter 1992; Baxter et al. 1992; Benkelfat et al. 1990; Swedo et al. 1992). Subsequent brain imaging studies have also implicated hyperfunction of the orbitofrontal cortex, basal ganglia, and the limbic system in OCD (Adler et al. 2000; Friedlander and Desrocher 2006; Kim et al. 2001; Mataix-Cols et al. 2004; McGuire et al. 1994; Menzies et al. 2008; Phillips and Mataix-Cols 2004; Rauch et al. 1994; Rotge et al. 2008; Saxena et al. 1998; van den Heuvel et al. 2004).

Deep brain stimulation (DBS) has been found effective for some patients with symptoms of OCD (Alegret et al. 2001; Anderson and Ahmed 2003; Fontaine et al. 2004; Mallet et al. 2002; Nuttin et al. 1999, 2003; Piasecki and Jefferson 2004), with the three sites targeted being the subthalamic nucleus (Fontaine et al. 2004; Mallet et al. 2002), the anterior limbs of the internal capsules (Anderson and Ahmed 2003; Nuttin et al. 2003), and the nucleus accumbens (Denys et al. 2010; Huff et al. 2010; Sturm et al. 2003). Although it is not well understood how DBS yields therapeutic effects, it may involve disruption of reverberatory activity in basal ganglia loops (Aouizerate et al. 2004; Bourne et al. 2012; Haynes and Mallet 2010; Tass et al. 2003), consistent with OCD pathophysiology. A similar disruption of overactivity may be expected from strategically placed lesions, and indeed severe cases of OCD may be improved with psychosurgery, in particular with anterior capsulotomy or cingulotomy (Baer et al. 1995; Chiocca and Martuza 1990; Dougherty et al. 2002; Jenike et al. 1991; Kettl and Marks 1986).

# 7 Neurotoxic Effects Producing OCD Pathophysiology

The pathophysiology of OCD – persistence of activity in SMS functional loops – could result from many types of insults and probably at several neuroanatomical locations in the neurocircuit. This section consults the clinical literature for medical conditions with a presentation of OCD phenomenology and asks whether those

medical perturbations could have yielded unregulated activity in basal ganglia functional loops. Three types of perturbations are considered: by drugs, brain injury, and infection.

## 7.1 Drugs

Given the prominent role of dopamine systems in SMS circuit (Fig. 2), it is expected that dopamine-activating drugs should provoke OCD phenomenology, at least with prolonged use. The clinical literature does indeed support this expectation with case reports of OCD symptoms from chronic exposure to amphetamines (Marchesi et al. 2009; Semple et al. 2011; Woolley and Heyman 2003), cocaine (Crum and Anthony 1993), and other direct dopamine receptor agonists (Ahlskog 2011). In animals, too, dopamine agonists produce compulsive behaviors considered to model OCD (De Carolis et al. 2011; Szechtman et al. 1999, 1998).

At first blush, a contradictory literature seemingly exists as well, because drugs used in treatment of OCD, dopamine receptors blockers (Bloch et al. 2006; Dougherty et al. 2004), may themselves induce OCD symptoms. There are clinical reports that atypical antipsychotics, such as aripiprazole (Desarkar et al. 2007), clozapine, risperidone, and olanzapine (Lykouras 2003), can produce or exacerbate OCD symptoms, as evidenced by the appearance and disappearance of symptoms with the introduction and removal of the drug. Clearly, those drug effects may involve an action on other than dopamine receptors. However, a possible mechanism via dopamine receptors should not be ignored, given that dopamine receptor supersensitivity may develop with exposure to dopamine blockers (Kostrzewa 1995) and given that dopamine D2 receptors play a role not only in the initiation of behavior but also in arresting ongoing activity (Ghahremani et al. 2012; Hoffman and Rueda Morales 2012). Indeed, the intricate circuitry of basal ganglia loops contains many inhibitory and excitatory connection points, not just dopaminergic or serotonergic (Bourdy and Barrot 2012; Pittenger et al. 2011; Rotge et al. 2010), and a disruption at any such point can yield overly sustained reverberatory activity.

## 7.2 Brain Injury

Symptoms of OCD often appear associated with neurologic disorders. These include postencephalitis Parkinsonian syndrome (Schilder 1938), Parkinson's disease (Kurlan 2004), striatal (Ercan et al. 2008) or globus pallidus necrosis (Laplane et al. 1989), Huntington's chorea (Cummings and Cunningham 1992), and Gilles de la Tourette's syndrome (TS) (Pauls et al. 1986). All of these conditions involve a disturbance of the basal ganglia, suggesting that upon activation of security motivation those neurologic disturbances may also yield the characteristic pathophysiology of OCD, namely, prolonged overactivity in the SMS neurocircuit. A similar inference of SMS circuit perturbation may relate to cases of OCD following acquired brain injury (Coetzer 2004).

## 7.3 Infection

Some instances of infection may result in OCD symptoms that resolve upon disappearance of the infection. For instance, there are such reported cases for cerebral malaria (Idro et al. 2005), *Mycoplasma pneumoniae* infection (Ercan et al. 2008), *Borna disease virus* (BDV) infection (Rotge et al. 2010), as well as streptococcal infection, the latter infection having a designation on its own for the ensuing OCD – PANDAS (pediatric autoimmune neuropsychiatric disorders associated with streptococcal infections) (Swedo et al. 1998, 1989). Some autoimmune disorders, including Sydenham's chorea (Swedo et al. 1989) and systemic lupus erythematosus (Yu et al. 2008), may also present with OCD symptoms. More generally, various immune markers may be altered in patients with OCD (da Rocha et al. 2008). Such findings suggest three types of mechanisms whereby infection and immune activation could yield OCD pathophysiology.

One type of mechanism follows the disease model of tuberculosis, namely, that a specific pathogen causes OCD, just like tubercle bacillus is the causal agent of tuberculosis. In tuberculosis, the symptoms and the pathophysiology of the disease are the outcomes of the bacterium lodging preferentially in certain body organs (e.g., lungs) and its interaction with immune processes to contain it. This battle yields "collateral damage" in the afflicted organs and physiology, and such particular perturbations are manifested as disease-specific symptoms. Similarly, for OCD, Rotge and colleagues (Rotge et al. 2010) elaborated a plausible scenario for "the process by which infection becomes disease" (Barnes 2000, p. 439). Following up on reports (Bode et al. 1995) that BDV infection may play a role in psychiatric disorders, Rotge and colleagues (Rotge et al. 2010) noted that the virus' preferential neurotropism for brain regions of relevance to OCD would mount an immune response there, leading possibly to a region-specific malfunction as a by-product of the immune response. While there are many plausible mechanisms by which immune reactions can perturb neuronal function (for a recent review, see Kapadia and Sakic (2011)), the authors highlight a mechanism which involves glutamate neuroexcitotoxicity, in light of genetic findings of a glutamate transporter gene SLC1A1 association with OCD (Arnold et al. 2006; Wu et al. 2012) and recognition that glutamate is both a neurotransmitter and an immunomodulator. According to the authors, genetic alterations of the glutamatergic system in some individuals would facilitate BDVinduced immunopathological reactions in the OCD-relevant brain regions (Rotge et al. 2010), with consequent hyperactivity of basal ganglia loops (Wu et al. 2012) that constitute the SMS neurocircuit.

Another type of mechanism is closely related to the above in that it also invokes immune activation as the agent producing neural dysfunction. However, this mechanism is related to autoimmune processes rather than stemming from local reactions to neurotropic pathogens. An example of such an autoimmuneinduced OCD-related neural dysfunction is PANDAS. The mechanism underlying the OCD of PANDAS is considered to involve an autoimmune response triggered by Group A beta-hemolytic streptococcal infection. In individuals positive for the D8/17 antigen on the surface of peripheral blood mononuclear cells, the antistreptococcal antibodies are thought to cross-react with neuronal basal ganglia antigens (Eisen et al. 2001; Swedo et al. 1997). This autoantibodies basal ganglia cross-reactivity can produce various forms of altered cellular function, including perturbation of neuronal signal transduction (Kapadia and Sakic 2011). Accordingly, OCD will ensue when humoral immunity targets neuronal elements of the SMS neurocircuit, and the resultant malfunction consists of overly persistent and uncontrolled circuit activity characteristic of OCD pathophysiology.

The last mechanism that can produce OCD pathophysiology is different: while still related to immune function, it does not involve an immunopathogenic effect on neuronal function. Rather, the OCD pathophysiology is produced by a sensitization-related mechanism that renders the SMS neurocircuit hyperfunctional. This may occur in the course of normal physiological function and not necessarily due to some immunopathogenic process. An example of such a phenomenon is "compensatory behavioral prophylaxis" (Fessler 2001). At times of relatively high progesterone levels, women are more susceptible to illness from infection because the immune system is suppressed, and a behavioral compensation at such times is enhanced precaution about contamination. Through a process of sensitization (Woolf 2011), this behavioral compensation mechanism can lead to SMS hyperactivity, producing OCD symptoms.

# 8 Conclusion

Because the cause or causes of OCD are unknown, it is uncertain whether variations in pathophysiology and symptoms constitute different manifestations of one disease or different diseases with distinct causes. However, whatever the causal agent may turn out to be, OCD is not a de novo creation but instead must reflect an exaggeration of the operation of a normal brain system. As argued here, information about the nature of the behavioral system affected by OCD can be integrated into a theoretical model of this normal system, called the security motivation system.

Recognizing both the similarities and differences between this system and other hard-wired motivational systems, a neurobiological model of the SMS may be derived from behavioral neuroscience studies of motivation. It consists of functional loops involving a cascade of cortico-striato-pallido-thalamo-cortical connections, with inhibitory connections from the brainstem. The pathophysiology of OCD can then be understood as normal neural activity in this system that becomes overly persistent and uncontrolled, e.g., due to a dopamine–serotonin imbalance. Three types of medical perturbations have been reported to produce OCD symptoms: drugs, brain injuries, and infections. The available evidence on these effects is broadly consistent with disturbances of regulation in the activity of the hypothesized basal ganglia functional loops. Thus, the SMS model provides a unifying framework for understanding the likely mechanisms of these neurotoxic effects.

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# Physical Exercise as Intervention in Parkinsonism

# Trevor Archer and Anders Fredriksson

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#### Abstract

Physical exercise, particularly for individuals with sedentary occupations and independent of activity type, offers probably the most effective health-ensuring policy available. The associations between physical exercise and symptoms/ biomarkers of idiopathic Parkinsonism and animal models of Parkinson's disease (PD), quality of life and self-reliance, disorder progression, and risk factors all support the contention that activity provides for an improved prognosis. In the present treatise, mice treated with the selective dopamine neurotoxin (1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine,  $3 \times 30$  mg/kg once each week over three weeks), or Vehicle, were given access to running-wheel exercise from the week

T. Archer (🖂)

Department of Psychology, University of Gothenburg, Gothenburg, Sweden e-mail: trevor.archer@psy.gu.se

A. Fredriksson

Department of Neuroscience, Psychiatry, Ulleråker, University Hospital, Uppsala, Sweden e-mail: anders.fredriksson@neuro.uu.se

following the first injection of MPTP onwards: the MPTP-Exercise group received four 30-min sessions in the running wheels (Mondays to Thursdays), whereas the MPTP No-exercise and Vehicle groups received a single session each week (Wednesdays). It was observed that the MPTP+Exercise group increased the distance run on each Wednesday 30-min session incrementally, whereas the MPTP No-exercise and Vehicle groups remained at the same distance throughout; similarly, during the 10-min test session on Fridays, prior to the tests of motor activity, the MPTP+Exercise group increased the distance run on each successive occasion but the MPTP No-exercise and Vehicle groups did not. In the tests of spontaneous motor activity, runningwheel exercise over 4 days/week (30 min, Mon.-Thurs.) improved all three parameters of motor activity, locomotion, rearing, and total activity, in the activity test chamber during test weeks 2-7, compared to the MPTP group following the MPTP administration. In the L-Dopa-induced activity test, running-wheel exercise over 4 days/week enhanced locomotor and rearing but not total activity following the subthreshold dose of L-Dopa for the MPTP+Exercise group but not the MPTP No-exercise group. Running-wheel exercise over 4 days/week increased the DA concentrations in the striatum of MPTP+Exercise mice.

#### Keywords

Alleviation • Biomarkers • Dopamine • L-Dopa • Mice • MPTP • Parkinson's disease • Physical exercise • Spontaneous motor activity • Symptoms • Wheel running

## 1 Introduction

Physical exercise has been described as any and all activity that generates force through muscular activity that disrupts a homeostatic state (McArdle et al. 1971; Scheuer and Tipton 1977). Although daily physical activity holds benefits for general measures of function, quality of life, and physical strength, as well as increasing endurance (Dechamps et al. 2010a, b; Marks et al. 2009), much evidence presents the manifest advantages for cerebral integrity during aging (Kramer et al. 1999; Lustig et al. 2009; Marks et al. 2010). Any bodily activity that enhances or maintains physical fitness implies the involvement of regular and frequent exercise. Morris and Schoo (2004) have defined exercise as a planned, structured physical activity with the purpose of improving one or more aspects of physical fitness and functional capacity. Regular physical exercise/exertion promotes neuroimmune functioning and facilitates prevention of heart conditions, cardiovascular diseases, type II diabetes and obesity, and mental health improvements, e.g., depressiveness, all of which may exacerbate the PD condition (Nicolucci et al. 2012; Stampfer et al. 2000). Physical exercise offers a nonpharmacologic, noninvasive intervention that enhances brain health and plasticity (Cotman and Berchtold 2002). It has been characterized on the basis of type, intensity, frequency, and duration, with either endurance or resistance as the training endpoint (Mougios 2010). Long-term exercise benefits brain functioning by increasing cerebral blood flow and oxygenation (Linkis et al. 1995), mobilizing growth factors and synaptic plasticity (Hunsberger et al. 2007) and facilitating performance through neurotransmitter release (Morishima et al. 2006; Hsu et al. 2011). Regular physical exercise holds particular benefits for older individuals, whether under conditions of normal aging or affected by neurodegenerative disorders (Archer 2011; Archer et al. 2011a, b).

Parkinson's disease (PD), a progressive brain disorder that impairs motor function, presents the cardinal symptoms of akinesia/hypokinesia, rigidity, and tremors as a result of the marked degeneration of the nigrostriatal dopamine (DA) pathway (Bernheimer et al. 1973; Goetz et al. 2001). This loss of DA in the basal ganglia is associated with deterioration in balance and postural control together with progressive reductions in the speed and amplitude of everyday movements (Morris et al. 2001; Smithson et al. 1998). The postural instability and disturbances in gait, which are major PD symptoms, are extremely disabling and lead to accidents, freezing, and involuntary movements, all of which worsen prognosis and elevate mortality risk (Adkin et al. 2003; Bloem et al. 2004; Wielinski et al. 2005). The progression of the disorder leads to reduced physical fitness and muscular capability (Canning et al. 1997) with psychosocial and economic implications for patients and caregivers (Schenkman et al. 2001). Fatigue presents a recurring symptom, described subjectively as "an overwhelming sense of tiredness, lack of energy, or feeling of exhaustion" (Krupp and Pollina 1996), that is linked to depressive symptoms and sleep disturbances with consequential loss of quality of life (Friedman and Friedman 2001; De Boer et al. 1996; Karlsen et al. 2000). Although there has been much advance in pharmacological and surgical interventions, postural instability, problems of gait, and instability of balance persist, thereby causing disablement (Grimbergen et al. 2004) and restriction of functional independence and quality of life (Martinez-Martin 1998; Schrag et al. 2000). Patients with PD may develop also a range of nonmotor conditions, including depression, apathy, sleep disturbances, constipation, and cognitive dysfunction, all of which together with the motor deficits may lead to adoption of a sedentary lifestyle (Fertl et al. 1993; van Nimwegen et al. 2011). Unsurprisingly, the prevalence of osteoporosis is high in patients with PD (63 % of PD women and 20 % of PD men compared with healthy age-matched control (29 % in women and 12 % in men) individuals (Invernizzi et al. 2009; Lorefält et al. 2007; Schuit et al. 2004), thereby increasing risk of falls (Sato et al. 2001).

It has been shown that physical exercise improves motor performance in PD patients (Bergen et al. 2002; Comella et al. 1994; Schenkman et al. 2008). In several controlled clinical studies, the implementation of continuous exercise for early-stage PD patients has provided improved daily activity, motor performance, ambulation, and overall functional independence (Baatile et al. 2000; Miyai et al. 2000; Reuter et al. 1999). In the laboratory, studies in healthy older rodents have shown that regular aerobic physical exercise promotes plasticity-related changes in the brain and CNS that include synaptogenesis neuronal arborization, enhanced

glucose utilization, angiogenesis, and neurogenesis (Hirsch and Farley 2009). It promotes also a healthy brain by reducing inflammation, suppressing oxidative stress, and stabilizing calcium homeostasis (Cotman et al. 2007). Several beneficial cerebral features of exercise have been observed from imaging studies: the volume and preservation of gray and white matters of sedentary individuals was enhanced through aerobic activity (Colcombe et al. 2006; Parker et al. 2011; Szabo et al. 2011). It has been suggested that in PD patients adaptive neuroplasticity may occur spontaneously at the level of both the basal ganglia and the cortex (Buhmann et al. 2005); these compensatory processes may be even initiated during the prodromal stage of disorder (van Nuenen et al. 2009). There is evidence that increased homocysteine (Hcy) levels might accelerate dopaminergic cell death in Parkinson's disease (PD) through neurotoxic effects (Gorgone et al. 2012: Lee et al. 2011; Zoccolella et al. 2010). Hyperhomocysteinemia presents a major risk factor for cerebral and peripheral vascular diseases as well as cortical and hippocampal injury, including an increased risk of dementia and cognitive impairment, and in PD patients treated with levodopa, elevated serum homocysteine (Hcy) concentrations are common. Nascimento et al. (2011) have found that even with regular levodopa therapy, Hcy concentrations in PD patients who had exercised regularly were significantly lower than in PD patients who had not exercised and were similar to the Hcy concentrations in the healthy controls. Taken together, evidence suggests that there is a strong tendency for PD patients toward sedentary lifestyle, whereas it is highly evident that a regime of regular physical exercise offers manifest benefits (Speelman et al. 2011; van Nimwegen et al. 2010).

Physical exercise is known to ameliorate both the symptoms and disability caused by PD (Bilowit 1956; Hurwitz 1989; Palmer et al. 1986). For example, the immediate effects of structured speed-dependent treadmill training, limited progressive treadmill training, conventional gait training, or a control intervention were compared in 17 early-stage PD patients. It was found that speed and stride length, the main gait disturbances, were improved markedly by speed-dependent training (Pohl et al. 2003). In another study, the effects of high-intensity training on the treadmill was compared with low- or no-intensity training with marked postexercise increases in gait speed, step, and stride by the former compared with the latter two conditions (Fisher et al. 2008). Treadmill physical activity has provided a variety of improvements for PD patients including motor performance (Schenkman et al. 2008), capacity for daily activities (Miyai et al. 2000, 2002), reduced incidence and lethality of disorder (Kuroda et al. 1992), strength, balance and gait speed (Toole et al. 2005), motor and postural flexibility (Kurtais et al. 2008), and neurocognitive performance (Ahlskog 2011; Gomez-Pinilla 2011). In addition to the cardinal symptoms, hypokinesia, gait and posture disturbance, tremors, and rigidity, several factors contribute negatively to reduced quality of life in PD patients: pain and fatigue, depressive symptoms and cognitive deficits, sleep problems, and detrimental social functioning with accompanying economical impact (Chrischilles et al. 2002; Whetten-Goldstein et al. 1997). Both functional mobility and qualityof-life variables have been improved by exercise programs of various different types (Herman et al. 2007; Kadivar et al. 2011; Rodrigues de Paula et al. 2006). Despite being slower compared to age-matched healthy controls, PD patients show motor learning with repetitive practice (Jessop et al. 2006; Michel et al. 2009; Onla-or and Winstein 2008; Smiley-Oyen et al. 2006). Both short-term and extensive practices of movement gave benefits against the bradykinesia of PD patients (Smiley-Oyen et al. 2012).

Repeated administration of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) to C57/BL6 mice induces selective and long-lasting lesions of DA in nigrostriatal regions of the brain (Jackson-Lewis et al. 1995; Jones-Humble et al. 1994). The neurotoxin, administered systemically, induces Parkinsonism in human and nonhuman primates (Langston 1985) that results in the loss of substantia nigra cells in the pars compacta of adult animals (Chiueh et al. 1985). It destroys selectively nigrostriatal neurons, thereby inducing acute, subacute, long-lasting, and even permanent effects that resemble certain features of PD, particularly the hypokinesic effects (Schultz et al. 1989). Systemic administration of MPTP ( $2 \times 40$ mg/kg, s.c.) caused L-Dopa reversible hypoactivity (Fredriksson et al. 1990; CR0105146 et al. 1990). A less rigorous dose regime, e.g.,  $2 \times 20$ , or 25 or 30 mg/kg, of MPTP has been found not to reduce motility in the C57 black mice, although DA brain concentrations may indicate up to 50-80 % reductions (Heikkila et al. 1989; Sonsalla and Heikkila 1986), unless given much more repeatedly (cf. Kurz et al. 2007). The parameters of MPTP treatment neurotoxicity in mice are extremely long-lasting (up to and beyond 52 weeks after treatment) with strong correlations between the functional deficits, particularly hypokinesia, the main biomarker, severe DA depletions, and a dose- and time-dependent recovery of several parameters of motor behavior following treatment with the DA precursor, L-Dopa (Archer and Fredriksson 2003; Fredriksson and Archer 1994; Fredriksson et al. 1999).

In the unilateral 6-hydroxydopamine rat model of Parkinson's disease (PD), Tillerson et al. (2003) abolished the lesion-induced motor asymmetry by forcing the rats to use affected (contralateral) limb, whereas as forced nonuse exacerbated the b). (DA) iniurv (Tillerson et al. 2002a. Both dopamine and 3.4dihydroxyphenylacetic acid (DOPAC) were elevated markedly in "casted" 6-OHDA-treated rats (forced to use the contralateral limb) compared with "noncasted" rats (Cohen et al. 2003). Archer and Fredriksson (2010) found that daily running-wheel activity attenuated the hypokinesic effects of MPTP in both a concentrated (2  $\times$  40 mg/kg, 24-h interval) and progressive (1  $\times$  40 mg/kg, weekly doses over 4 weeks) schedule with regard to spontaneous motor behavior and activity following a subthreshold dose of L-Dopa. The loss of DA in each case was attenuated by exercise also (Experiment I, 61 % of control rather than 17 %, Experiment II, 24 % rather than 11 %). Using the progressive schedule of MPTP treatment and extending the exercise intervention from 7 to 14 weeks, it was shown that spontaneous motor activity after MPTP was close to restoration, whereas activity after subthreshold L-Dopa was completely recovered (Fredriksson et al. 2011); DA levels were restored from 17 % (non-exercised) to 64 % in the 14-week exercise intervention and levels of brain-derived neurotrophic factor were increased significantly. It was shown also that both the functional and DA deficits

by MPTP were attenuated even by delayed- introduction of exercise (Archer and Fredriksson 2012).

Brain-derived neurotrophic factor (BDNF) is a neurotrophin with widespread expression in the brain and is connected intimately with brain metabolism and homeostasis (Chaldakov 2011). It is associated with neurogenesis, neuronal survival, and neuroreparation in the brain and CNS (Cui 2006; Numakawa et al. 2010). Treatment interventions that enhance BDNF-related signaling have the potential to restore neural connectivity (Kaplan et al. 2010). Physical exercise induces improvements in motor ability and enhances BDNF expression (Macias et al. 2009). It is linked to elevated BDNF levels in the hippocampus (Neeper et al. 1996; Oliff et al. 1998). Voluntary running, as physical activity, amplifies the BDNF signal that augments neurogenesis through diverse molecular pathways (Stranahan et al. 2009). BDNF mediates several essential morphological changes at neuronal levels that include dendritic arborization (Imamura and Greer 2009; Zhou et al. 2008). axonal and dendritic remodeling (Menna et al. 2005), synaptogenesis (Liu et al. 2009; Tchantchou et al. 2009), and synaptic efficacy (Boulanger and Poo 1999; Sallert et al. 2009). Faherty et al. (2005) have shown that a combination of exercise, social interactions, and learning, or exercise alone during adulthood, gave total protection against MPTP-induced Parkinsonism. They found also that changes in mRNA expression suggested that increases in glia-derived neurotrophic factors, coupled with a decrease of dopamine-related transporters (e.g., dopamine transporter, DAT; vesicular monoamine transporter, VMAT2), contributed to the observed neuroprotection of dopamine neurons in the nigrostriatal system following MPTP exposure. Tajiri et al. (2010) observed that exercise induced behavioral recovery in an animal model of PD and caused increased BDNF and glial-derived neurotrophic factor (GDNF) in the striatum of 6-OHDA-treated rats.

In the pathophysiology of PD symptom profiles, not least with regard to the cardinal symptoms and fatigue, L-Dopa treatment, despite certain drawbacks, has been found to improve ventilator function (Vincken et al. 1989) and work endurance and efficiency without affecting exercise hemodynamics (LeWitt et al. 1994). As DA precursor, the compound, following acute administration, reduced cortisol concentrations in PD patients in resting positions that reduced stress conditions (Müller and Muhlack 2007; Müller et al. 2006); this effect has been related to alterations in serotonergic and dopaminergic transmission (Höglund et al. 2001; Khakimova et al. 2011; Lopez et al. 2001). Pickrell et al. (2011) have presented a mouse model that expresses a mitochondria-targeted restriction enzyme, PstI or mito-PstI, damaging mitochondrial DNA (mtDNA) in DA neurons by inducing double-strand breaks in the mtDNA, leading to an oxidative phosphorylation deficiency, mostly due to mtDNA depletion. Using a dopamine transporter (DAT) promoter-driven tetracycline transactivator protein (tTA), they expressed mito-PstI exclusively in DA neurons, thereby presenting a novel PD transgenic mouse model (PD-mito-PstI mouse). They have shown that the PD-mito-PstI mice present the major features of PD: possessing a motor phenotype that is reversible with L-Dopa treatment, a progressive neurodegeneration of the SN dopaminergic population, and striatal DA depletion. Their results showed also that behavioral phenotypes in PD-mito-PstI mice were associated with striatal dysfunctions preceding SN loss of tyrosine hydroxylase-positive neurons and that other neurotransmitter systems [noradrenaline (NE) and serotonin (5-HT)] were increased after the disruption of DA neurons, potentially as a compensatory mechanism. The eventual effects of regular physical exercise upon the PD-mito-PstI mice are awaited. The administration of L-Dopa, despite development of tolerance, generally alleviates bradykinesia (Fredriksson and Archer 1994; Fredriksson et al. 1990, 1999), but clinically, the DA precursor by itself will not elevate movement unless assertive physical exertion in the early stages is provoked (Weiner and Singer 1989). It has been found that exercise induced a significantly faster stimulation of growth hormone release following L-Dopa release, in comparison with the rest condition, in patients with PD (Müller et al. 2007).

Although endurance training aerobic sessions enhance cortical release proportional to exercise intensity (Kanaley et al. 2001), intensive physical exercise sessions also reduced cortisol plasma concentrations (Kemmler et al. 2003). Forced exercise initially increased, and then reduced, adrenocorticotropic hormone (ACTH) in control, non-brain damaged animals, followed by marked increases in BDNF (Griesbach et al. 2012). Exercise effects upon Parkinsonian conditions are complex: it appears that PD patients treated with L-Dopa may report a decreased capacity for physical exercise (Reuter et al. 2000; Ziv et al. 1998). The influence of physical exercise as an intervention affecting the efficacy of L-Dopa upon Parkinson symptoms is a central issue: Müller and Muhlack (2008) reported that while endurance exercise did not affect the reduction in cortisol release by L-Dopa, grip strength increased only during rest. Although exercise seems not to affect the pharmacokinetics of L-Dopa in PD patients (Carter et al. 1992), there is an abundance of evidence showing that it impacts upon L-Dopa effects on motor parameters in PD (Muhlack et al. 2007; Reuter et al. 1999; Scandalis et al. 2001; Viliani et al. 1999). Frazzitta et al. (2012) have found that intensive rehabilitation treatment, that included treadmill and stabilometric platform training, reduced the daily medication dosage of L-Dopa in treated (exercised) patients, whereas it was significantly increased in control (non-exercised) patients. In addition, the beneficial effects of intensive rehabilitation treatment persisted over time with a second rehabilitation cycle administered after one year as effective as the initial treatment. Endurance exercise has the effect of increasing L-Dopa-induced growth hormone availability in PD patients (Keranen et al. 1996; Manetta et al. 2002): Müller et al. (2007) observed that during exercise the resulting higher blood pressure mobilized blood flow and thereby growth hormone-releasing hormone, secreted into the hypothalamic portal capillaries, toward the growth hormone-producing cells in the pituitary gland. In highlighting the unique importance of exercise in the disorder perspective, a recent review (Ahlskog 2011) made the following recommendations for clinicians: (1) ongoing vigorous exercise and physical fitness should be encouraged, (2) PD physical therapy programs ought to include structured, graduated fitness instruction and guidance for deconditioned patients with PD, and (3) levodopa and other forms of dopamine replenishment therapy should be utilized to achieve the maximum capability and motivation whereby patients maintain fitness.

The purpose of the present study was to examine whether or not physical exercise in a running wheel would ameliorate the functional and DA deficits induced by MPTP under a progressive dose regime of the neurotoxin.

# 2 Material and Methods

## 2.1 Animals

Male C57 Bl/6 mice were purchased from B&K, Sollentuna, Sweden, and were maintained, five to a cage, in plastic cages in a room at temperature of  $22 \pm 1$  °C and a 12/12 h constant light/dark cycle (lights on between 06.00 and 18.00 h). They were placed and maintained in groups of 4–6 animals in a room maintained for male mice only following arrival at the laboratory for about 2 weeks in order to acclimatize. Free access to food and water was maintained throughout, except for the day previous to the initiation to running-wheel exercise which occurred at the end of the second week following arrival. They were housed in groups of six animals, running-wheel exercised and activity chamber tested only during the hours of light (08.00–15.00 h). All exercising and testing were performed in a normally lighted room. Half of the mice in the MPTP treatment condition (MPTP-Exercise) were given running-wheel exercise, whereas the other half were placed in a clean laboratory cage for the same period in a room in which the running wheels were placed. Motor activity was tested in a specially arranged test room. This test room, in which all 12 ADEA activity test chambers, each identical to the home cage, were placed, was well secluded and used only for this purpose. Each test chamber (i.e., motor activity test cage) was placed in a sound-proofed wooden box with 12 cm thick walls and front panels and a small double-glass window to allow observation; each box had a dimmed lighting.

Three weeks following arrival, three groups (n = 10) of DSP4-treated and two groups of vehicle-treated mice were administered either MPTP ( $2 \times 40$  mg/kg, s.c., 24 h between injections) or vehicle (0.9 % physiological saline injected s.c. in a volume of 2 ml/kg body weight). Milmed (see below for details of preparation) or vehicle were administered on one occasion per week (Experiment I) or twice weekly (Experiments II and III).

Experiments were carried out in accordance with the European Communities Council Directive of 24 November 1986 (86/609/EEC) after approval from the local ethical committee (Uppsala University and Agricultural Research Council) and by the Swedish Committee for Ethical Experiments on Laboratory Animals (license S93/92 and S77/94, Stockholm, Sweden).

# 2.2 Drugs

MPTP (Research Biomedical Inc., MA, USA;  $2 \times 20$  mg/kg or  $2 \times 40$  mg/kg, s.c., with a 24-h interval between injections in each case) was dissolved in saline and

administered s.c. in a volume of 2 ml/kg body weight. Melmid .... Saline was used as vehicle in each case.

# 3 Behavioral Measurements and Apparatus

Activity test chambers: An automated device, consisting of Makrolon rodent test cages ( $40 \times 25 \times 15$  cm), each placed within two series of infrared beams (at two different heights, one low and one high, 2 and 8 cm, respectively, above the surface of the sawdust, 1 cm deep), was used to measure spontaneous motor activity (RAT-O-MATIC, ADEA Elektronik AB, Uppsala, Sweden). The distance between the infrared beams was as follows: the low-level beams were 73 mm apart lengthwise and 58 mm apart breadthwise in relation to the test chamber and the high-level beams, placed only along each long side of the test chamber, were 28 mm apart. According to the procedures described previously (Archer et al. 1986), the following parameters were measured: LOCOMOTION was measured by the low grid of infrared beams. Counts were registered only when the mouse is in the horizontal plane, ambulating around the test cage. REARING was registered throughout the time when at least one high-level beam was interrupted, i.e., the number of counts registered was proportional to the amount of time spent rearing. TOTAL ACTIVITY was measured by a sensor (a pickup similar to a gramophone needle, mounted on a lever with a counterweight) with which the test cage was constantly in contact. The sensor registered all types of vibration received from the test cage, such as those produced both by locomotion and rearing as well as shaking, tremors, scratching, and grooming. All three behavioral parameters were measured over three consecutive 20-min periods. The motor activity test room, in which all 12 ADEA activity test chambers, each identical to the home cage, were placed, was well secluded and used only for this purpose. Each test chamber (i.e., activity cage) was placed in a sound-proofed wooden box with 12 cm thick walls and front panels, and daylighting. Motor activity parameters were tested on one occasion only, over three consecutive 20-min periods, at the age of 3–4 months.

# 4 Design and Treatment

On Friday, week 1, mice were injected single weekly injections of MPTP ( $1 \times 30 \text{ mg/}$ kg, s.c.) or vehicle after a test of spontaneous motor activity, without access to wheel-running activity prior to the first administration of MPTP. Access to wheel-running activity was given on 1 day only for the Vehicle and MPTP groups (Wednesdays, 30 min) but on 4 days each week for the MPTP+Exercise (Mondays to Thursdays, 30 min), and this 1-day versus 4-day exercise regime was continued up to and during the 7th week. On Friday, weeks 2 and 3, mice were injected single weekly injections of MPTP ( $1 \times 30 \text{ mg/kg}$ , s.c.) or vehicle after a test of spontaneous motor activity. In each case, each mouse in each of the four groups received a 10-min access to the running wheel just prior to the 60-min spontaneous motor activity test (Table 1).

	Vehicle	MPTP	MPTP+Exer		
Monday	Cage	Cage	Cage		
	Cage	Cage	Cage		
Week 1	Exer	Exer	Exer		
	Cage	Cage	Cage		
Friday: T🛛	Test+sal	Test+MPTP	Test+MPTP		
Måndag	Cage	Cage	Exer		
	Cage	Cage	Exer		
Weeks 2–3	Exer	Exer	Exer		
	Cage	Cage	Exer		
Fri: T2&30	Test+Sal	Test+MPTP	Test+MPTP		
Måndag	Cage	Cage	Exer		
	Cage	Cage	Exer		
Weeks 4–7	Exer	Exer	Exer		
	Cage	Cage	Exer		
Fri. T4,5,6,70	Test	Test	Test		
Fri.T*	Test	Test	Test		
Week 8	Sacrifice and neu	Sacrifice and neurochemical analysis			

**Table 1** The experimental design and treatment of mice administered either MPTP or Vehicle, with access to running wheel exercise either 1 day or 4 days each week

To SMA only; T\* SMA+L-Dopa

Vehicle = Saline (0.9% koksalt-lösning), Mptp = **30 mg/kg** OBS!!! subcutan injektion av 5 ml/kg Exer = 30 minuter i löphjul, Cage = 30 minuter i ensam bur Y = Yeast 10 ml/kg S = Saline 10 ml/kg, Test = Spontan aktivitet under  $3 \times 20$  minuter, Inducerad aktivitet efter injektion av L-Dopa (5 mg/kg i en volym av 5 ml/kg)

# 4.1 Neurochemical Analysis

## 4.1.1 Dopamine

Mice were killed by cervical dislocation within 1–2 weeks of completion of behavioral testing. Determination of DA was performed using a high-performance liquid chromatography with electrochemical detection (HPLC-EC), according to Björk et al. (1991), as modified by Liu et al. (1995). Striatal regions were rapidly dissected out and stored at -80 °C until neurochemical analysis. DA concentration was measured as follows: the frozen tissue samples were weighed and homogenized in 1 ml of 0.1 M perchloric acid, and alpha-methyl-5-hydroxytryptophan was added as an internal standard. After centrifugation (12,000 rpm, i.e., 18,600 g, 4 °C, 10 min) and filtration, 20  $\mu$ l of the supernatant was injected into the HPLC-EC to assay DA. The HPLC system consisted of a PM-48 pump (Bioanalytical Systems, BAS) with a CMA/240 autoinjector (injection volume: 20  $\mu$ l), a precolumn (15 × 3.2 mm, RP-18 Newguard, 7  $\mu$ m), a column (100 × 4.6 mm, SPHERI-5, RP-18, 5  $\mu$ m), and an amperometric detector (LC-4B, BAS, equipped with an Ag/AgCl reference electrode and a MF-2000 cell) operating at a potential of +0.85V. The mobile phase, ph 2.69, consisted of K2HPO4 and citric acid buffer (pH 2.5),

10 % methanol, sodium octyl sulfate, 40 mg/l, and EDTA. The flow rate was 1 ml/min, and the temperature of the mobile phase was 35  $^{\circ}$ C.

## 4.2 Statistical Analysis

The locomotion, rearing, and total activity data over three consecutive 20-min periods in the activity test chambers from the spontaneous motor activity data were submitted to a split-plot ANOVA design (Kirk 1995). Brain (striatal) regional levels of dopamine, locomotion, rearing, and total activity over the full 60-min period following administration of apomorphine, each were submitted to a one-way ANOVA based on a completely randomized design (Kirk 1995). Pairwise testing between the different treatment groups was performed with the Tukey's HSD test (Kirk 1995). The 1 % level of significance was maintained throughout unless where otherwise stated.

#### 4.2.1 Results

#### Distance/30 min: Wednesday Tests

Neither the MPTP nor the vehicle groups showed any increase in distance run over 30 min in the running wheels, whereas the MPTP+Exercise group increased significantly distance run over 30 min from test week 2 to test week 7. The former received only one 30-min exposure (Wed.) per week to the wheels, while the latter received four 30-min exposures (Mon.–Thurs.) per week. Split-plot ANOVA indicated a significant Groups x test week interaction effect F(20, 251) = 25.83, p < 0.0001, for distance during 30 min. Figure 1 presents the distance run by each group during 30 min in the running wheels on the Wednesday tests.

Pairwise testing with Tukey's HSD test indicated significant differences between the three groups over test weeks 2–7 and a significant increase by the MPTP+Exercise group over test weeks 2–7.

#### Distance/10 min: Friday Tests

Neither the MPTP nor the vehicle groups showed any increase in distance run over 10 min in the running wheels, whereas the MPTP+Exercise group increased significantly distance run over 10 min from test week 2 to test week 7. Split-plot ANOVA indicated a significant Groups x test week interaction effect F(20, 251) = 47.36, p < 0.0001, for distance during 10 min. Figure 2 presents the distance run by each group during 10 min in the running wheels on the Friday tests.

Pairwise testing with Tukey's HSD test indicated significant differences between the three groups over test weeks 2–7 and a significant increase by the MPTP+Exercise group over test weeks 2–7.

#### Spontaneous Motor Activity: Friday Tests

Running-wheel exercise over 4 days/week (30 min, Mon.–Thurs.) improved all three parameters of motor activity: locomotion, rearing, and total activity, in the activity test chamber during test weeks 2–7, compared to the MPTP group

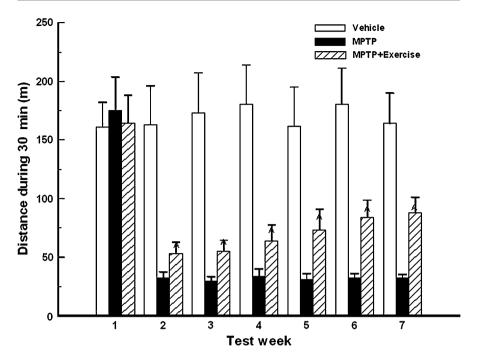


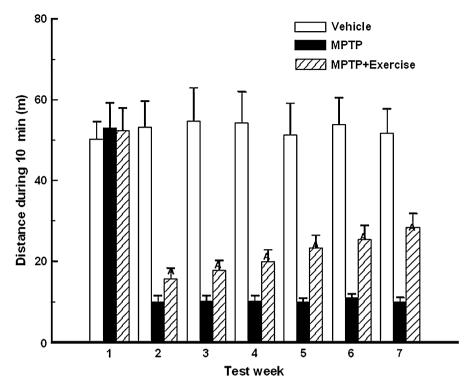
Fig. 1 Distance covered during the 30-min Wednesday test by the Vehicle, MPTP, and MPTP+Exercise groups

following the MPTP administration. Thus, split-plot ANOVA indicated a significant Groups x test week interaction effect for locomotion, rearing, and total activity: F(20, 209) = 18.82, p < 0.0001; F(20, 2309) = 15.36, p < 0.001; and F(20, 209) = 7.44, p < 0.0001, respectively. Figure 3 presents the locomotion, rearing, and total activity by each group during 60 min in the activity test chambers on the Friday tests.

Pairwise testing with Tukey's HSD test indicated significant differences between the three groups over test weeks 2–7 and a significant increase by the MPTP+Exercise group over test weeks 2–7.

#### L-Dopa-Induced Activity: Friday Tests

Running-wheel exercise over 4 days/week enhanced locomotor and rearing but not total activity following the subthreshold dose of L-Dopa for the MPTP-Exercise group but not the MPTP No-exercise group. Thus, one-way ANOVA indicated a significant Groups effect for locomotion, rearing, and total activity: F(2, 23) = 39.95, p < 0.0001; F(2, 23) = 81.21, p < 0.0001; and F(2, 23) = 4.77, p < 0.02, respectively. Figure 4 presents the locomotion, rearing, and total activity by each group during 180 min in the activity test chambers, following L-Dopa (5 mg/kg) on the Friday tests.



**Fig. 2** Distance covered during the 10-min Friday test by the Vehicle, MPTP, and MPTP+Exercise groups

Pairwise testing with Tukey's HSD test indicated significant differences between the three groups over test weeks 2–7 and a significant increase by the MPTP+Exercise group over test weeks 2–7.

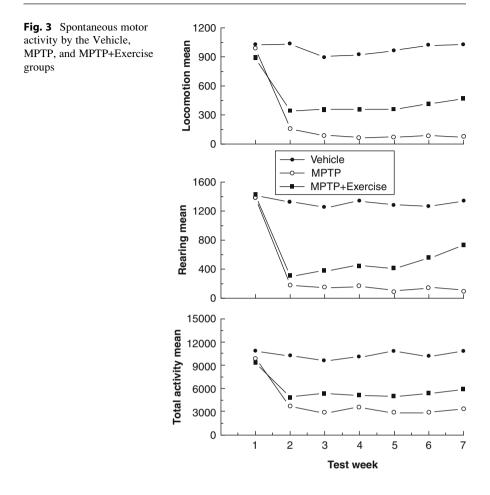
#### **Dopamine Analysis**

Running-wheel exercise over 4 days/week increased the DA concentrations in the striatum of MPTP+Exercise mice. Thus, one-way ANOVA indicated a significant Groups effect for DA concentration: F(2, 17) = 121.03, p<0.0001. Figure 5 presents the DA concentrations in the striatum of the Vehicle, MPTP, and MPTP+Exercise groups.

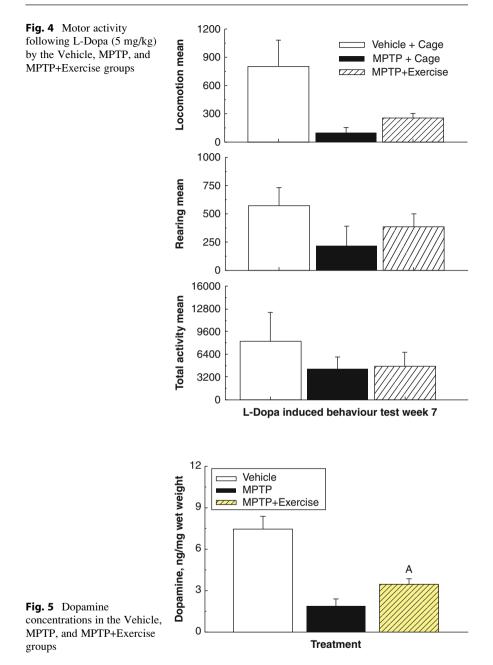
Percent of Vehicle control analyses indicated that striatal DA in the MPTP group was 11 % compared with 31 % in the MPTP-Exercise group.

# 5 Discussion

The present account of the utility of physical exercise (daily wheel-running activity, 4 days/week as opposed to 1 day/week) in alleviating the functional and



neurochemical deficits induced by MPTP may be summarized as follows: (1) Only the MPTP+Exercise group, not the Vehicle and MPTP groups, showed significant increases in distance run over 30 min from test week 2 to test week 7 on the Wednesday tests. (2) Only the MPTP+Exercise group, not the Vehicle and MPTP groups, showed significant increases in distance run over 10 min from test week 2 to test week 7 on the Friday tests, given prior to the tests in the motor activity test chambers. (3) The tests of spontaneous motor activity indicated that running-wheel exercise over 4 days/week (30 min, Mon.–Thurs.) improved all three parameters of motor activity, locomotion, rearing, and total activity, in the activity test chamber during test weeks 2–7 (on Fridays), compared to the MPTP group following the MPTP administration. (4) The L-Dopa-induced activity tests indicated that runningwheel exercise over 4 days/week enhanced locomotor and rearing but not total activity following the subthreshold dose of L-Dopa administered on Fridays after the spontaneous motor activity tests. (5) Running-wheel exercise over 4 days/week



increased the DA concentrations in the striatum of MPTP+Exercise mice compared with the MPTP (one-day/week exposure to the running wheels only) group.

The present findings confirm and extend out previous observations (Archer and Fredriksson 2010, 2012; Fredriksson et al. 2011) in several respects. As observed

previously, physical exercise (30 min/day, 4 days/weeks) reduced the hypokinesic effects of MPTP and the DA loss induced by the neurotoxin under various conditions and parameters. However, none of the previous studies incorporated any method for establishing the exact distance run during each exercise bout by each mouse in its running wheel at any determined time interval, i.e., how much did each day of exercise. The present study incorporated also the condition that "non-exercised" groups in fact received one 30-min wheel-running exposure per week (Wednesdays) such that the exercise variable in fact involved one (non-exercise condition) as opposed to four (exercise condition). As the results (see Figs. 1 and 2, particularly) indicate, a single 30-min exercise session/week does nothing to affect exercise propensity, whereas four sessions/week produced a limited, but significant, and steady increase in distance run.

In the Archer and Fredriksson (2010) study, DA analyses indicated that after 4  $\times$ 40 mg/kg MPTP accompanied by 5 weeks of exercise (four 30-min/day sessions per week), the MPTP no-exercise group indicated 11 % of Vehicle control values. whereas the MPTP-Exercise group indicated 24 % of control values. In Experiment II of the Fredriksson et al. (2011) study applying the same  $4 \times 40$  mg/kg MPTP administration regime but accompanied by a 4-week exercise schedule (four 30-min/ day sessions per week), DA analyses indicated that the MPTP no-exercise showed 17 % of vehicle controls, whereas the MPTP-Exercise showed 64% of the vehicle controls. Taken together, a comparison of the results from these two studies suggests that longer periods of weekly exercise promote improved recovery from the MPTP-induced DA depletions. In the Archer and Fredriksson (2012) study, both experiments employed a  $4 \times 30$  mg/kg MPTP administration regime accompanied by an 8-week exercise schedule (four 30-min/day sessions per week) in both Experiments I and II. For both experiments, exercise was introduced only after the second injection of MPTP. It was observed that DA levels in Experiment I were 20 % of control value for the MPTP no-exercise group and 39 % of control values for the MPTP-Exercise group; in Experiment II, DA concentrations were 23 % and 43% for the MPTP no-exercise and MPTP-Exercise groups, respectively. In the present experiment, a  $3 \times 30$  mg/kg MPTP administration regime, with exercise introduced after the 1st MPTP injection (except for Wednesday) accompanied by a 6-week exercise schedule (four 30-min/day sessions per week). DA concentrations were 11 % of control values for the MPTP no-exercise group and 31 % for the MPTP-Exercise group. A plausible consensus from these MPTP experiments, employing single, weekly doses of MPTP, appears to be that independent of MPTP dose level or number of injections, the introduction of a multiple weekly exercise regime, rather than a single weekly session regime, significantly restores DA concentrations in the striatum as well as some restoration of motor activity.

The recovery from loss of DA presents an essential aspect of physical activitymediated interventions in animal models of Parkinsonism and in particular those applying MPTP dose regimes. Petzinger et al. (2007) administered a series of 4 injections of MPTP (i.p.), or saline at 2-h intervals for a total of 80 mg/kg; physical exercise consisted of treadmill running sessions on an accelerating rotarod was initiated for half of the MPTP and half of the saline mice 5 days later. All the mice that were provided access to running-wheel exercise showed increased latencies to fall off the accelerating rotarod, thereby demonstrating an improved balance, compared with the non-exercised mice. Kurz et al. (2007) administered male C57/BL6 mice with 10 injections of MPTP (25 mg/kg) and probenecid (250 mg/kg) over 5 weeks; the control group of mice received probenecid only, and all the mice were monitored on a motorized treadmill. They observed that the MPTP-treated mice showed significantly more variable stride length and less certain gait pattern than the control mice. In the Ahmad et al. (2009) study, the effects of physical exercise using a motorized rodent treadmill (speeds up to 15 m/min, 40 min/day, 5 days/week for 10 and 18 weeks) for mice given chronic treatment with MPTP (12.5 mg/kg  $\times$  10, with a 3.5-day interval between doses, and in combination with probenecid, 250 mg/kg) were examined. This chronic MPTP treatment regime induces a 52 % loss of neurons in the ventral tegmental area (VTA), including decrease in cell volume and irregular or disparaging axonal and dendritic projections that express deficits in dendritic arborization morphology in animals that remain sedentary (German et al. 1996). Positron emission tomography (PET)-imaging studies applying DA-D2R radiotracers have provided the prospect of performing longitudinal studies of the effects of exercise in both humans and animals. Nevertheless, no alteration in DA release or alteration in the binding of [11C]raclopride as a result of aerobic exercise was observed although DA release was triggered (Wang et al. 2000). Vuckovic et al. (2010) found that in MPTP mice high-intensity treadmill exercise, using PET imaging with [<sup>18</sup>F]fallypride, induced an increase in striatal DA-D2R expression.

In PD patients treated with L-Dopa, elevated serum homocysteine (Hcy) concentrations are commonly observed (Müller 2011: Müller and Muhlack 2010; Sui and Zhang 2010). Nascimento et al. (2011) have showed that Hcy concentrations in PD patients who exercised regularly were significantly lower than in sedentary patients but were similar in healthy controls. Regular exercise improves other aspects of brain health and integrity: for example, endurance exercise promoted cardiorespiratory rehabilitation in MPTP mice with severe neurodegeneration (Al-Jarrah et al. 2007). PD results in significant decreases in blood vessel density (Geny et al. 1994). Al-Jarrah et al. (2010) showed that four weeks of treadmill exercise training induced an increased striatal blood vessel density in chronic Parkinsonian mice that had received 10 administrations of MPTP (25 mg/kg) and probenecid (250 mg/kg) over 5 weeks. They found also that exercise induced an increase in striatal blood vessel density in the control mice although this effect did not attain significance. Smith et al. (2011) measured recovery of gait performance and amount of spontaneous physical activity using a "parallel rod activity chamber" (PRAC) in MPTP-treated and control mice provided with treadmill running (1h/day at 18 cm/s, 5 days/week) or not. They observed that the treadmill exercise program improved gait performance and increased physical activity as well as promoting increased protein expression of striatal dopamine transporter and tyrosine hydroxylase in both MPTP and control mice, but most markedly in the former.

Also applying the chronic MPTP mouse model of PD with moderate neurodegeneration, Lau et al. (2011) observed that exercise improved mitochondrial function and produced increases in region-specific levels of brain-derived and glial cell line-derived neurotrophic factors (BDNF and GDNF). They have demonstrated the impact of exercise upon protection against nigrostriatal neuronal, mitochondrial, and locomotor deficits in the moderate chronic mouse model of PD. Tajiri et al. (2010) examined the effects of compulsive exercise on Parkinson's disease model of rats. Bromodeoxyuridine (BrdU) was injected to label proliferating cells prior to 6-hydroxydopamine (6-OHDA, 20 µg) lesion into the right striatum of female SD rats. Later, at 24 h after the lesion, the rats were forced to run on the treadmill (5 days/week, 30 min/day, 11 m/min). For behavioral analysis, cylinder tests were performed at 1, 2, 3, and 4 weeks and amphetamine-induced rotational test was performed at 2 and 4 weeks followed by sacrifice for immunohistochemical investigations. The exercise group showed better behavioral recovery in cylinder test and significant decrease in the number of amphetamine-induced rotations, compared to the non-exercise group. Correspondingly, significant preservation of tyrosine hydroxylase (TH)-positive fibers in the striatum and TH-positive neurons in the substantia nigra pars compacta (SNc) was demonstrated, compared to the nonexercise group. In addition, the number of migrated BrdU- and doublecortin-positive cells toward the lesioned striatum was increased in the exercise group. BDNF and GDNF were increased in the striatum by exercise. Their observations suggest that exercise exerts neuroprotective effects or enhances the neuronal differentiation in Parkinson's disease model of rats with subsequent improvement in deteriorated motor function. Finally, Patki and Lau (2011) have demonstrated that long-term exercise (over 18 weeks) attenuated both cytochrome c release and elevated levels of p53; both proteins are associated with mitochondrial dysfunction in the chronic MPTP mouse model. Cytochrome c is released by the mitochondria in response to proapoptotic stimuli. The p53 pathway responds to stresses that can disrupt the fidelity of DNA replication and cell division. A stress signal is transmitted to the p53 protein by posttranslational modifications that causes the activation of the p53 protein as a transcription factor that initiates a program of cell cycle arrest, cellular senescence, or apoptosis (e.g., Culmsee and Mattson 2005; Mihara et al. 2003).

#### 6 Conclusion

Taken together, the present findings and the accumulation of laboratory and human studies reinforce the contention that physical exercise may be highly beneficial in conditions of mild to moderate Parkinsonism.

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# Protective Agents in Parkinson's Disease: Caffeine and Adenosine A2A Receptor Antagonists

# Nicola Simola, Annalisa Pinna, Lucia Frau, and Micaela Morelli

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## Abstract

The pharmacologic management of Parkinson's disease is based on drugs that act on the motor symptoms, whereas there are currently no drugs available that can alter the progressive neurodegeneration of dopaminergic neurons. Based on recent findings suggesting that the adenosinergic system is one of the most interesting in the field of neuroprotection in Parkinson's disease, this chapter describes the functions of adenosine and its receptors in the central nervous system, with particular emphasis on their role in neurotoxicity/neuroprotection. Results of epidemiologic surveys demonstrating that intake of caffeine, an adenosine  $A_1/A_{2A}$  receptor antagonist, is inversely correlated with Parkinson's

N. Simola • L. Frau

A. Pinna • M. Morelli (⊠)

Department of Biomedical Sciences, University of Cagliari, Cagliari, Italy e-mail: nicola.simola@gmail.com; luciafrau79@hotmail.com

Department of Biomedical Sciences, University of Cagliari, Cagliari, Italy

CNR Institute of Neuroscience, Cagliari, Italy e-mail: apinna@unica.it; morelli@unica.it

disease are summarized. Moreover, evidence originating from preclinical studies showing that the antagonism of the adenosine  $A_{2A}$  receptor is responsible for the neuroprotective effects of caffeine is also presented. This chapter therefore provides a comprehensive analysis of the current literature concerning the adenosinergic-based neuroprotective intervention strategy for Parkinson's disease.

# 1 Introduction

Parkinson's disease (PD) is the second most prevalent neurodegenerative disease, with an occurrence that tends to rise dramatically with increasing age, thereby amplifying the social relevance of the disease as life expectancy increases (de Rijk et al. 2000). The degeneration of dopaminergic neurons of the substantia nigra pars compacta (SNc) that project to the striatum, the hallmark of PD, provokes functional modifications within the basal ganglia (BG) circuitry causing the typical motor symptoms (tremor, rigidity, and bradykinesia). PD is a classic movement disorder; however, the motor symptoms, which appear when the dopaminergic neuron degeneration has reached at least 70–80 %, are preceded by numerous non-motor symptoms, olfactory deficits, and gastrointestinal and cognitive dysfunctions (Poewe 2008). Idiopathic PD is a complex disorder characterized by the involvement of selected neuronal populations throughout the central and peripheral nervous systems; however, neurotransmitters other than dopamine (DA) are involved in the disease, including noradrenaline, serotonin, and acetylcholine (Jellinger 1991).

The primary cause of the degenerative process underlying PD remains unknown; however, several contributing factors have so far been identified for the idiopathic form of the disease. The most important of these include mitochondrial defects, oxidative damage, anomalous protein aggregation, and neuroinflammation (Schapira 2006). These processes, once initiated, continue to produce dopaminergic neuron liability contributing to the loss of efficacy of DA-replacement therapy.

The main therapeutic developments in the field of PD have focused, first, on improvement of DA-replacement therapies in order to better manage or prevent the onset of motor complications and, second, on the discovery of compounds that could modify the course of neurodegeneration. Among the various therapeutic approaches aimed at addressing these objectives, the manipulation of adenosine neurotransmission is one of the most valuable.

The first suggestion that manipulating adenosine neurotransmission might have a beneficial effect on PD onset or progression came from epidemiologic evidence showing that consumption of caffeine, a nonselective  $A_1/A_{2A}$  receptor antagonist, reduced the risk of developing PD (Ascherio et al. 2001; Costa et al. 2010) (see Sect. 5). Furthermore, preclinical studies investigating the adenosine receptor type involved in these effects suggested that the blockade of the  $A_{2A}$  receptor subtype provides the best neuroprotective effect (Schwarzschild et al. 2006).

# 2 Neuromodulatory Role of Adenosine

Adenosine is an endogenous purine nucleoside constitutively present in mammalian tissues, where it plays a regulatory role in a variety of important physiologic processes by acting as a homeostatic modulator. Adenosine is produced as a result of hydrolysis of adenosine monophosphate (AMP) by means of an action of ecto-5'-nucleotidase (Fredholm et al. 2005); thus, its formation depends upon the metabolism of adenosine triphosphate (ATP). In the extracellular compartment, the level of adenosine also depends on the rate of hydrolysis of ATP, which is released from either neurons or glial cells, and on the adenosine carrier, which keeps adenosine at a stable level. The actions of adenosine are mediated by specific G-protein-coupled receptors. To date, four adenosine receptors have been cloned and characterized:  $A_1$ ,  $A_{2A}$ ,  $A_{2B}$ , and  $A_3$  (Fredholm et al. 2005).

Adenosine plays different roles in normal physiology, which include promoting and/or maintaining sleep, regulating the general state of arousal, and coupling cerebral blood flow to energy demand (Dunwiddie and Masino 2001). Many of the effects of adenosine that are observed in normal conditions are modified during pathologic events, and the multiple roles of adenosine may produce dual neuroprotective/neurotoxic effects based on the type of insult and cellular condition (Cunha 2001).

Endogenous adenosine released during hypoxia, ischemia, electrical activity, hypoglycemia, or aglycemia reduces the subsequent damage to the neuronal tissue. The neuroprotection offered by adenosine is also effective against other kinds of damage that are not as directly related to energy metabolism, such as mechanical cell injury (Mitchell et al. 1995), methamphetamine and 3,4-methylenedioxymethamphetamine (MDMA)-induced neuroinflammation, and neurotoxicity (Delle Donne and Sonsalla 1994; Khairnar et al. 2010). In these conditions, the neuroprotective actions of adenosine are mediated primarily via  $A_1$  receptor activation, whereas it seems that the action on the  $A_{2A}$  receptor could result in a neurotoxic effect. Caffeine and other  $A_{2A}$  antagonists, in fact, have induced neuroprotection and reduced glial cell activation in animal models of dopaminergic neurotoxicity utilizing 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) or 6-hydroxydopamine (6-OHDA) (Schwarzschild et al. 2006; Carta et al. 2009; Frau et al. 2011).

#### 3 Adenosine A<sub>2A</sub> Receptors

Among the different adenosine receptors, of particular interest for PD are the  $A_{2A}$  receptors, specifically located in the BG circuit (Fig. 1). The adenosine  $A_{2A}$  receptors interact functionally and are co-expressed with dopaminergic  $D_2$  receptors on the striatal GABAergic neurons of the "indirect" BG pathway projecting from the striatum to the globus pallidus (Hettinger et al. 2001). In the BG circuit,  $A_{2A}$  receptors are prevalently localized in the axons, soma, dendrites, and dendritic

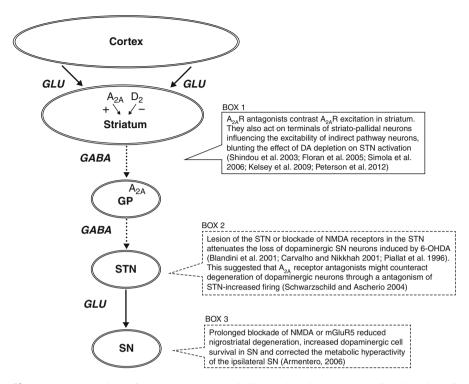


Fig. 1 Proposed actions of  $A_{2A}$  receptors on the indirect striatonigral pathway. Classical view of BG functions demonstrating that dopaminergic neuron degeneration provokes an increase in glutamatergic input from the cortex to the striatum and an increase in GABAergic indirect output from the striatum to the GP, leading to an increase in STN activity. In turn, an increase in STN glutamatergic activity contributes to excitotoxic dopaminergic neuron degeneration in the SN. As shown in Box 1, several reports indicate that  $A_{2A}$  receptors control excitability of the striatopallidal GABAergic pathway and that A2A receptor antagonism counteracts the effects of dopaminergic neuron degeneration. As a result, the GABAergic control from the GP to the STN is increased, leading to a decrease in STN activity. As shown in Box 2, a decrease in STN activity and glutamatergic output from the STN to the SN counteracts excitotoxic degeneration of the dopaminergic neurons in the SN. As shown in Box 3, blockade of glutamatergic receptors in the SN reduces excitotoxic degeneration of dopaminergic neurons. Therefore, one of the mechanisms of neuroprotection by A<sub>2A</sub> receptor antagonists may be the indirect inhibition of STN activity. Box 1 (continuous line) represents a direct action of A2A receptor antagonist; Boxes 2 and 3 (dashed line) represent an indirect action of A<sub>2A</sub> receptor antagonist on neurotransmitter release. Abbreviations: BG basal ganglia, DA dopamine, GLU glutamate, GP globus pallidus, mGluR5 metabotropic glutamate receptor 5, SN substantia nigra, STN subthalamic nucleus, 6-OHDA 6-hydroxydopamine

spines of the striatal GABAergic neurons of the indirect pathway. In contrast, very few striatonigral neurons of the "direct" pathway, containing dopaminergic  $D_1$  receptors, express  $A_{2A}$  receptors (Schiffmann and Vanderhaeghen 1993) (Fig. 1). However, activation or blockade of  $A_{2A}$  receptors in the indirect striatopallidal pathway impairs or facilitates dopaminergic  $D_1$ -mediated responses as well (Ferre et al. 1997).

In light of the neuromodulatory role of adenosine,  $A_{2A}$  receptors have been described to interact directly or indirectly with several receptors, such as dopaminergic  $D_2$  and  $D_3$  receptors, NMDA, metabotropic glutamate receptor types 4 (mGLUR<sub>4</sub>) and 5 (mGLUR<sub>5</sub>), cannabinoid receptor type 1 (CB1), and 5-hydroxytryptamine receptor type 1A (5-HT<sub>1A</sub>), and to form heteromeric complexes with some of them (Kurokawa et al. 1996; Gerevich et al. 2002; Łukasiewicz et al. 2007; Armentero et al. 2006, 2011; Bogenpohl et al. 2012; Jones et al. 2012).

Besides the presence of  $A_{2A}$  receptors in neurons, their existence has been described in astrocytes: specifically, about 3 % of  $A_{2A}$  receptors were found in glial cells of the striatum (Hettinger et al. 2001).

## 4 Caffeine Properties

Caffeine is a natural xanthine alkaloid, contained in several popular dietary sources, such as coffee, tea, chocolate, and soft drinks. Its ability to elicit psychostimulatory effects, that are usually not accompanied by harmful unwanted consequences, renders caffeine the most consumed psychoactive substance (Fredholm et al. 1999).

At doses in the range of normal consumption, caffeine induces its effects by blocking the central adenosine  $A_1$  and  $A_{2A}$  receptors (Fredholm et al. 1999). Caffeine-induced psychostimulant effects appear to be influenced by the dopaminergic transmission since enhancement of caffeine-elicited psychomotor activation is observed in experimental rodents following administration of dopaminergic agonists (Cauli and Morelli 2005), whereas an attenuation of the effects of caffeine is present after administration of dopaminergic antagonists (Green and Schenk 2002). The influence that caffeine exerts on dopaminergic transmission is of great interest, considering the critical involvement of DA in the modulation of important functions, such as movement, attention, goal-directed behavior, and associative learning (Robbins and Roberts 2007; Phillips et al. 2008).

The existence of negative correlations between PD and dietary factors, such as caffeine, is of particular interest, in the light of caffeine's safety and widespread consumption (see Sect. 5). In this regard, the results from preclinical studies demonstrating that caffeine can exert protective effects in different models of neurotoxicity add further importance to the finding (Table 1).

Caffeine administration has been shown to have a neuroprotective effect in rodent and primate models of dopaminergic neuron degeneration, such as the MPTP and 6-OHDA paradigms (Table 1). Caffeine neuroprotection is particularly important since it is afforded by a substance with negligible toxic or adverse effects. Often, substances proposed to be neuroprotective impair neuronal functions essential for correct cellular homeostasis or alter the levels of neurotransmitters, such as glutamate, which are important for cognitive functions. Caffeine, in contrast, is a safe substance used worldwide with no unwanted effects in the broad population.

Table 1 Overview	Table 1 Overview of the most common paradigms of toxin-induced neurodegeneration used to investigate neuronal damage and neuroprotection in PD	oxin-induced neurodegeneration	used to investigate neuronal damage a	ind neuroprotection in PD
Model	Type	Toxic effects observed	Main features of the model	Effect of caffeine administered before or concomitantly to the toxin
Cell cultures	Acute Can use different cells (e.g., neuroblastoma or mesencephalic cells) and toxins (e.g., 6-OHDA, MPP <sup>+</sup> , rotenone)	Decrease in cell viability	Allows to rapidly identify putative neuroprotective agents and to study their molecular mechanisms Can be used as a complement of in vivo models	Cytoprotection <sup>4: b</sup>
6-OHDA in rats	Acute Employs the direct intracerebral infusion of the toxin	Degeneration of DA neurons in the SNc Reduction in the striatal content of DA and its metabolites	The degeneration is rapid and reaches a near terminal stage few days after infusion Oxidative stress is the major mechanism of neurotoxicity Allows to selectively target different brain regions according to the site of toxin infusion	Attenuation in the demise of SNc neurons <sup>c</sup> Restoration of the levels of DA and its metabolites in the striatum <sup>c</sup>
Lipopolysaccharide in rats and mice	Acute or chronic Loss of DA neurons in t Can use either intracerebral infusion Activation of glial cells or systemic administration Elevation in the levels c neuroinflammatory med	Loss of DA neurons in the SNc Activation of glial cells Elevation in the levels of neuroinflammatory mediators	Induces a selective degeneration of DA neurons in SNc, while sparing other DA neurons Degeneration of DA neurons is delayed, progressive, and irreversible Neuroinflammation is the main mechanism underlying neuronal death	Attenuation of neuroinflammation <sup>d</sup>
Paraquat or Paraquat + maneb in mice	Chronic	Degeneration of DA neurons in the SNc Loss of DA terminals in the striatum	This model may yield variable results Attenuation in the and significant mortality degeneration of D. The toxins used are clinically relevant, since they have been suggested as a potential causative factor of PD	Attenuation in the degeneration of DA neurons <sup>1</sup>

MPTP in mice	Acute or acute-repeated with multiple administration at a fixed interval The toxin is administered systemically	Degeneration of DA neurons in the SNc Loss of DA terminals in the striatum Reduction in the striatal levels of DA and its metabolites DA and its metabolites Glial activation	The severity of DA lesion can be adjusted by varying the protocol of administration It is still questioned whether the lesion is progressive The model can be used to study both neurotoxicity and neuroinflammation The toxin acts chiefly on mitochondrial function The toxin is clinically relevant, as MPTP can induce PD in humans	Attenuation in the degeneration of SNc degeneration of SNc neurons <sup>e.</sup> Restoration of the levels of DA and its metabolites in the striatum <sup>6,1,g</sup> Counteraction of neuroinflammation <sup>h</sup>
Rotenone in rats and mice	Chronic The toxin is administered by osmotic minipumps or parenteral injection	Degeneration of SNc DA neurons Loss of striatal DA terminals accompanied by reduction in striatal DA content	DA neuron degeneration is often accompanied by other nonspecific brain damages Systemic toxicity and high mortality is often associated with, or precedes, neurodegeneration There is a high variability in the results obtained with the use of this model Differently from other commonly used models, chronic rotenone can generate proteinaceous inclusions in DA neurons resembling those observed in human PD The toxin used is clinically relevant, since exposure to rotenone has been suggested as a possible causative factor of PD	Not tested
The table describes some of model are reported, together Abbreviations: <i>DA</i> dopamii substantia nigra pars compao <sup>a</sup> Nakaso et al. 2008; <sup>b</sup> Nobre- 2009; <sup>i</sup> Kachroo et al. 2010	The table describes some of the most common toxin-based experimental models of parkinsonian-like model are reported, together with the effects of caffeine on the degenerative process, when available Abbreviations: $DA$ dopamine, $MPP^+$ 1-methyl-4-phenylpyridinium, $MPTP$ 1-methyl-4-phenyl-1,2, substantia nigra pars compacta, 6-0HDA 6-hydroxydopamine <sup>a</sup> Nakaso et al. 2008; <sup>b</sup> Nobre et al. 2010; <sup>c</sup> Aguiar et al. 2006; <sup>d</sup> Brothers et al. 2010; <sup>c</sup> Schwarzschild et al 2009; <sup>†</sup> Kachroo et al. 2010	ed experimental models of parkin on the degenerative process, when Ipyridinium, <i>MPTP</i> 1-methyl-4-1 mine J6; <sup>d</sup> Brothers et al. 2010; <sup>e</sup> Schwarz	The table describes some of the most common toxin-based experimental models of parkinsonian-like neuron degeneration. The main characteristics of each model are reported, together with the effects of caffeine on the degenerative process, when available Abbreviations: <i>DA</i> dopamine, <i>MPP</i> <sup>+</sup> 1-methyl-4-phenylpyridinium, <i>MPTP</i> 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine, <i>PD</i> Parkinson's disease, <i>SNc</i> substantia nigra pars compacta, 6-OHDA 6-hydroxydopamine al. 2010; <sup>e</sup> Schwarzschild et al. 2003; <sup>b</sup> Nobre et al. 2010; <sup>c</sup> Aguiar et al. 2006; <sup>d</sup> Brothers et al. 2010; <sup>e</sup> Schwarzschild et al. 2003a; <sup>f</sup> Chen et al. 2001; <sup>g</sup> Xu et al. 2010; <sup>h</sup> Carta et al.	: main characteristics of each D Parkinson's disease, <i>SNc</i> <sup>8</sup> Xu et al. 2010; <sup>h</sup> Carta et al.

# 5 Epidemiologic Studies on the Neuroprotective Role of Caffeine in Parkinson's Disease

Numerous findings suggest that the two decades before the manifestation of motor impairments in PD are fundamental in the onset of the progressive demise of dopaminergic neurons. Therefore, epidemiologic studies investigating habits, health status, and lifestyle of individuals who developed PD are very important in order to obtain new information about PD development and to identify new approaches to prevent neurodegeneration in PD.

One of the most interesting results that has emerged from these epidemiologic studies is the inverse correlation between caffeine intake and risk of developing PD. The three main epidemiologic studies, that provided consistent suggestions regarding this relationship, are the Honolulu Heart Program (HHP, 30 years of follow-up of  $\sim 8.000$  Japanese-American men), the Health Professionals Follow-up Study (HPFS, 10 years of follow-up of ~47,000 men), and the Nurses' Health Study (NHS, 16 years of follow-up of ~88,500 women) (Ross et al. 2000; Ascherio et al. 2001, 2003). In these surveys, besides information of lifestyles and health status of individuals, nutritional behaviors, including data about consumption of coffee and/ or other caffeinated or decaffeinated beverages, were collected. The results of the investigation demonstrated that individuals consuming medium to high quantities of caffeine have a decreased risk of developing PD (Ross et al. 2000; Ascherio et al. 2001, 2003). Importantly, this finding was dose dependent and was adjusted for age, smoking status, alcohol use, other nutrients, or potential confounding variables (Ascherio et al. 2001, 2003). The studies described that individuals consuming caffeine (including tea and all caffeine sources) had a lower risk of developing PD compared with non-regular caffeine drinkers (less than one cup a day), whereas no association was found with consumption of decaffeinated coffee. Moreover, a 50 %risk reduction of PD was observed among men consuming less than one cup a day compared with male non-coffee drinkers (Ross et al. 2000; Ascherio et al. 2001). Notably, this inverse relationship between caffeine intake and PD risk was not so evident in women (Ascherio et al. 2001, 2003). Indeed, a clear gender difference was demonstrated by Ascherio and coworkers (2001), regarding the association between caffeine intake and risk of PD in women in the NHS. Similar findings have been reported by Benedetti and coworkers (2000). Since gender difference with regard to the effect of caffeine suggested a possible hormonal cause, in the women reported in the NHS, the correlation between use of postmenopausal estrogens, caffeine intake, and risk of PD was examined. Generally, use of postmenopausal estrogens was not correlated with risk of developing PD. On the other hand, among estrogens users, women consuming more than five cups of coffee per day were found to have a higher risk of PD compared with women who never drink coffee. In contrast, similar to men, women who were coffee drinkers and had never taken postmenopausal estrogens had a lower risk of PD than non-coffee drinkers (Ascherio et al. 2001). These data were confirmed by two separate prospective studies involving men and women which provided evidence that caffeine may decrease risk of PD only in men and in women who did not take postmenopausal estrogens, but not among estrogen consumers (Ascherio et al. 2003, 2004). Furthermore, the potential beneficial effect of caffeine might be prevented by estrogen therapy (Ascherio et al. 2003, 2004). Conversely, two prospective studies of Finnish individuals reported no gender differences in the opposite correlation between coffee intake and PD risk (Hu et al. 2007; Sääksjärvi et al. 2008). However, in the study by Saaksjarvi and coworkers (2008), the percentage of women was only about 5 % of the cohorts; therefore, the effect of estrogen therapy could not be examined for this small number of women.

Recently, an experimental study in the MPTP mouse model of PD investigated the biologic basis of the interaction between estrogen and caffeine consumption on risk of PD. This study showed that caffeine attenuated, dose dependently, the MPTP-induced striatal DA loss in both male and ovariectomized female compared with female non-ovariectomized mice (Xu et al. 2006). Moreover, chronic estrogen treatment prevented the neuroprotection induced by caffeine in male and in female ovariectomized mice, confirming that the hormonal therapy might prevent the neuroprotective effect of caffeine in this model of PD (Xu et al. 2006). Metabolic or pharmacokinetic interactions between caffeine and estrogens have been supposed, but it is not clear whether this interaction is fundamental in the reduced neuroprotective effect observed when estrogen therapy and caffeine are combined (Xu et al. 2006).

A recent statistical study by Costa and coworkers (2010) collected and statistically analyzed the majority of epidemiologic studies, which better evaluate the effect of caffeine intake and the incidence of PD, up to September 2009. This statistical review, which includes 26 studies, unequivocally confirms the inverse association between caffeine exposure and the risk of developing PD (Costa et al. 2010).

In spite of the epidemiologic and experimental findings strongly supporting the concept that dietary caffeine decreases the possibility of developing PD, very little is known about the correlation between caffeine intake and the degree of disease progression in PD patients. Two clinical studies, in which the progression of the disease and the consumption of several caffeinate beverages were evaluated by the modified Unified Parkinson's Disease Rating Scale (UPDRS), did not demonstrate clear evidence of the neuroprotective role of caffeine with respect to degree of PD progression (Schwarzschild et al. 2003b; Simon et al. 2008). Therefore, further larger investigations are required to fully assess the neuroprotective role of caffeine on disease progression in PD patients.

# 6 Mechanisms of Dopaminergic Neuroprotection by Caffeine and A<sub>2A</sub> Receptors Blockade

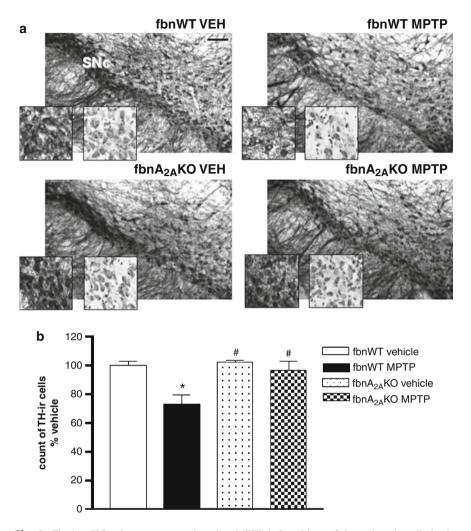
Studies of neuroprotection in PD should carefully consider a number of factors with regard to the features of neurodegeneration, as well as to the experimental model in which the study is performed. In recent years, PD has been increasingly envisaged as a multifactorial pathology, and several factors have been proposed to be involved in the degeneration of dopaminergic neurons (Schapira 2006). Therefore, it is

hypothesized that neuroprotective drugs should act by means of multiple mechanisms rather than a single mechanism. Moreover, it is worth mentioning that most of the experimental models of PD in use employ toxins according to an acute, or acute-repeated, protocol of administration (Table 1). This may significantly differ from idiopathic PD, with respect to time course of degeneration and mechanisms involved in neuronal demise. Therefore, the choice of experimental model is a critical step when addressing the mechanisms of neuroprotective agents.

Several studies in rodent models of PD demonstrate that  $A_{2A}$  receptor antagonists counteract both the demise of dopaminergic neurons in the SNc and the drop of DA levels in the striatum (Schwarzschild et al. 2003a; Morelli et al. 2010). However,  $A_{2A}$  antagonists exert beneficial effects not only in models of PD neurodegeneration but also in paradigms of Alzheimer's and Huntington's disease-like neurotoxicity (Stone et al. 2009). This evidence suggests that  $A_{2A}$  receptor antagonists counteract PD-like neurodegeneration by means of mechanisms which are not selective towards dopaminergic neurons but are part of a broader neuroprotective action. In the light of these observations, the two major hypotheses proposed to explain the neuroprotective effects of  $A_{2A}$  antagonists on dopaminergic neurons concern the modulation of glutamate-induced excitotoxicity and neuroinflammation, two mechanisms involved in several neurodegenerative pathologies.

Glutamate-mediated excitotoxicity has long been envisaged as an important player in PD neurodegeneration. Thus, dopaminergic neurons are vulnerable to changes in glutamate extracellular concentrations, on the one hand, while a dysregulation of glutamate transmission has been described in PD, on the other (Lancelot and Beal 1998; Greenamyre 2001). Remarkably, the stimulation of adenosine  $A_{2A}$  receptors can elevate the extracellular concentrations of glutamate, while blockade of  $A_{2A}$  receptors decreases glutamate extracellular levels which, in turn, may relieve the excitotoxic insult and afford neuroprotection towards dopaminergic neurons (Popoli et al. 1995). Recent evidence supports a role for nonneuronal cells in the modulation of glutamate release by  $A_{2A}$  receptor blockade (Melani et al. 2003; Yu et al. 2008), which is in line with the localization of these receptors on glial cells (see Sect. 3).

Several reports have shown that  $A_{2A}$  receptor antagonists modulate the activity of the indirect pathway by acting on the terminals of striatopallidal medium spiny neurons or by influencing the intrinsic excitability of indirect pathway spiny projection neurons, blunting the effects of DA depletion (Shindou et al. 2003; Floran et al. 2005; Simola et al. 2006; Kelsey et al 2009; Peterson et al. 2012) (Fig. 1). Moreover, independent investigations have shown that both lesion of the subthalamic nucleus (STN) and blockade of the NMDA receptors in the STN attenuate the loss of dopaminergic neurons induced by 6-OHDA (Piallat et al. 1996; Blandini et al. 2001; Carvalho and Nikkhah 2001). Based on these considerations, release of glutamate from the STN can be envisioned as an important mechanism in dopaminergic neuron degeneration. Moreover, it has to be considered that the overactivation of the STN progresses in the course of PD, and this can eventually exacerbate the loss of dopaminergic neurons over time. Therefore, it is



**Fig. 2** FbnA<sub>2A</sub> KO mice are protected against MPTP-induced loss of dopaminergic cells in the SNc. (**a**) Representative sections from SNc immunostained for TH. Insets show higher magnification of TH-labeled (*left*) and cresyl violet-labeled (*right*) cells. Mice were treated with MPTP (20 mg/kg once a day for 4 days) or vehicle. (**b**) Graph shows analysis of TH immunostaining in fbnA<sub>2A</sub> KO mice, reported as a percentage of TH-positive cells compared with vehicle-treated mice. \*p < 0.05 versus WT vehicle-treated group; #p < 0.05 versus WT MPTP-treated group, by Tukey's post hoc test. Scale bar, 50 µm. Abbreviations: *Fbn* forebrain neurons, *KO* knock-out, *MPTP* 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine, *SNc* substantia nigra pars compacta, *TH* tyrosine hydroxylase, *WT* wild type. (Reproduced from Carta et al. 2009)

conceivable that  $A_{2A}$  antagonists, by regulating the activity of the striatopallidal pathway, besides counteracting motor disability, may also attenuate degeneration of the dopaminergic neurons through a modulation of STN firing (Schwarzschild and Ascherio 2004) (Fig. 1).

An important mechanism involved in the neurodegenerative process that characterizes PD is neuroinflammation, as suggested by preclinical and epidemiologic evidence (Hirsch and Hunot 2009; Halliday and Stevens 2011). Neuroinflammation gives rise to a complex cascade of events in the brain, resulting in the activation of glial cells (microglia, astrocytes, and oligodendrocytes) and the subsequent generation of inflammatory mediators (e.g., cytokines, interferons, interleukins) (Reale et al. 2009). Even though transient neuroinflammation can be beneficial, as it may help the brain in coping with noxious insults, protracted neuroinflammation is detrimental and may lead to neuronal damage. It has been recently suggested that glial activation plays an important role in the initiation of early response and in the progression of degenerative diseases, such as PD, in which  $\alpha$ -synuclein accumulation in astrocytes causes recruitment of phagocytic microglia that attack selected neurons causing neuronal degeneration (Halliday and Stevens 2011). Furthermore, not only can neuroinflammation promote neurodegeneration, but neuronal death itself can trigger inflammatory responses, which may aggravate the ongoing neuronal damage (Litteljohn et al. 2010).

Both microglia and astrocytes express  $A_{2A}$  receptors, the stimulation of which promotes the recruitment of these cells in neuroinflammatory responses (Lopes et al. 2011). Conversely, blockade of  $A_{2A}$  receptors suppresses the activation of microglia and astrocytes, as well as the stimulation of the proinflammatory signal cascade (Armentero et al. 2011). Interestingly, recent evidence obtained in the mouse MPTP model of PD demonstrates that the increased survival of dopaminergic neurons mediated by  $A_{2A}$  antagonists is accompanied by a significant attenuation in microgliosis and astrogliosis (Yu et al. 2008; Carta et al. 2009; Frau et al. 2011).

Based on these premises and on the current knowledge of  $A_{2A}$  antagonists and neuroprotection, it is feasible that attenuation of glutamate-mediated excitotoxicity and neuroinflammation may participate in the protective effects of  $A_{2A}$  antagonists on dopaminergic neurons. However, it is important to consider that in experimental models of PD, adenosine  $A_{2A}$  receptor antagonists counteract the degeneration of dopaminergic neurons at doses lower than those required to elicit motor stimulation (Yu et al. 2008; Carta et al. 2009). It is therefore hypothesized that excitotoxicity and neuroinflammation are phenomena clearly distinct from regulation of motor function, as for the mechanisms involved. Furthermore, excitotoxicity and neuroinflammation have been envisaged as causative factors in neurodegenerative disorders other than PD (Glass et al. 2010). Therefore, an attenuation, or regulation, of these noxious phenomena by  $A_{2A}$  antagonists, besides justifying the protective effect observed in PD models, would be in agreement with the data demonstrating that these drugs can protect neurons from a wide array of insults.

Even though the hypotheses on excitotoxicity and neuroinflammation are intriguing, caution should be taken when addressing the mechanisms underlying the neuroprotection mediated by  $A_{2A}$  antagonists in models of PD. In this connection, it is interesting to point out that astrocytes, the activation of which has been found to be increased in the MPTP model of PD (Carta et al. 2009), regulate not only neuroinflammation but also glutamate extracellular levels and, accordingly, excitotoxicity (Melani et al. 2003). Therefore, astrocytes could be a crucial point at

which  $A_{2A}$  antagonists may promote the survival of dopaminergic neurons, perhaps by attenuating both excitotoxicity and neuroinflammation. Even though the evidence supporting the involvement of glial elements in the dopaminergic neuroprotection by  $A_{2A}$  antagonists is increasingly growing, neuronal mechanisms cannot be disregarded. Thus, recent evidence demonstrates that the inactivation of neuronal forebrain A2A receptors completely counteracts the degeneration of dopaminergic neurons in the MPTP model of PD (Carta et al. 2009) (Fig. 2). Furthermore, it has to be considered that a number of factors have been proposed to promote and/or sustain the degeneration of dopaminergic neurons in PD, besides excitotoxicity and neuroinflammation. These include, for example, oxidative stress, proteolytic damage, and mitochondrial defects (Schapira 2006). Therefore, it is conceivable that several different mechanisms may participate in the rescue of dopaminergic transmission by  $A_{2A}$  antagonists. In this connection, it is interesting to note that  $A_{2A}$  antagonists have been reported to protect dopaminergic neurons in a model of mitochondrial dysfunction (Alfinito et al. 2003). In addition, it has been suggested that the different causative factors of PD neurotoxicity may engender a sort of vicious cycle, with some triggering initial neuronal damage and others perpetuating it until neuronal death is reached. Therefore,  $A_{2A}$  antagonists could intervene at different steps of this cycle and, by interrupting it, afford neuroprotection.

To date, blockade of adenosine  $A_{2A}$  receptors has been demonstrated to efficiently restore motor function in PD, without exacerbating dyskinesia, both in animal models and in clinical trials of PD patients (Hauser and Schwarzschild 2005; Schwarzschild et al. 2006; Morelli et al. 2010). Therefore, their role as disease-modifying drugs in PD is of specific relevance since  $A_{2A}$  antagonists may unequivocally affect PD with regard to both symptoms and neuroprotection.

#### 7 Conclusion

Caffeine and adenosine  $A_{2A}$  receptor antagonists have given new impetus to the research on PD in the latest 10 years. In the absence of effective neuroprotective treatments for PD, epidemiologic studies investigating dietary factors, such as caffeine, that may allow individuals to lower the risk of PD became compelling. Remarkably, preclinical studies have shown that the neuroprotective potential of caffeine extends to chronic administration of pesticides (Kachroo et al. 2010), suggesting that caffeine can prevent degeneration of dopaminergic neurons induced by a broad range of agents, including environmental toxins, in addition to pathophysiologic mechanisms.

Moreover, the preclinical and clinical investigations suggesting that  $A_{2A}$  receptor antagonists represent a class of drugs that might counteract motor deficits symptomatically and, at the same time, delay or halt the progression of dopaminergic neuron degeneration are very important for future delineation of effective therapies for this disease. It should, however, be underlined that the

discovery of neuroprotective strategies in chronic neurodegenerative diseases, such as PD, largely depends on the possibility of monitoring the progression of the neurodegeneration. The availability of biomarkers associated with disease progression therefore becomes crucial for the advancement of this strategy. Although the research on biomarkers is very promising (Halperin et al. 2009), none of them can predict PD with 100 % confidence nor provide a clear definition of subgroups at risk. Therefore, the future of research in PD clearly depends on the possibility of developing reliable and affordable biomarkers in order to facilitate and render the clinical trials of new molecules consistent.

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# Stem Cell Therapies for Age Associated Neurodegeneration

# Stephanie Merchant, Sarah Stegeman, and Kiminobu Sugaya

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## Abstract

Regenerative medicine is the newly emerging field of this decade, namely, stem cell therapeutics. Interestingly, the field of stem cell therapeutics has been around for decades, such as the treatment of Leukemia using bone marrow transplantations; however, little momentum has been generated in the field until recently. During this predominantly stationary period of the regenerative movement, many issues have been debated regarding stem cell therapeutics. Many questions to these past issues are now starting to rise to the forefront due to breakthrough research being conducted in the regenerative field: adult versus embryonic stem cell populations, ethics, ease of access to these populations, and, more importantly, what types of diseased states can these stem cell populations be effective in? In order to answer this last question, it is imperative to

S. Merchant (⊠) • S. Stegeman • K. Sugaya

College of Medicine, University of Central Florida, Orlando, FL, USA e-mail: stephanie.brucker@gmail.com; ksugaya@me.com; ksugaya@mail.ucf.edu

understand not only basic stem cell pathologies but also the pathological conditions of the diseased state to be treated. Perhaps one of the most challenging to overcome is Alzheimer's disease (AD), a neurodegenerative condition characterized by a predominant loss of adult stem cell populations. What makes AD a challenge is the need to overcome both the natural effects of aging as well as the caustic environmental conditions. In order to introduce new neuronal networking through stem cell therapeutics, a better understanding of the pathological environment is needed. With new and upcoming technology, such as induced pluripotent stem cells, used to produce neural cells from somatic cells, and small molecular compounds which increase endogenous stem cell populations, it will be possible to perform autologous regenerative therapies, thus providing potential cures for age-related diseases such as AD.

List of Abbrevia	
AD	Alzheimer's disease
APOE4	Apolipoprotein E-e4
APP	Amyloid precursor protein
AS	Adult stem cells
β-amyloid	Beta amyloid
BACE	Beta-site APP-cleaving enzyme
BBB	Blood–brain barrier
bFGF	Basic fibroblast growth factor
BMPs	Bone morphogenetic protein
CCg	Glycosylated form of cysteine C
CDC	Center for Disease Control
ChEIs	Cholinesterase inhibitors
CNS	Central nervous system
CNTF	Ciliary neurotrophic factor
DNA	Deoxyribonucleic acid
DS	Down syndrome
EGF	Epidermal growth factor
eNS	Endogenous neural stem cells
ES	Embryonic stem
FDA	Food and Drug Administration
GFAP	Glial fibrillary acidic protein
gp130	Glycoprotein 130
hNSC	Human neural stem cells
IGF-1	Insulin-like growth factor
IL-6	Interluekin-6
iPS	Induced pluripotent stem
JAK	Janus kinase
Klf4	Krüppel-like family of transcription factors
LIF	Leukemia inhibitory factor
MAPK	Mitogen-activated protein kinase
MS	Mesenchymal stem

NFT	Intracellular neurofibrillary tangles
Ngn-1, Ngn-2	Neurogenin 1 and 2
NP	Neural progenitor
NS	Neural stem
Oct 3/4	Octamer-binding transcription factor 3/4
PS-1, PS-2	Presenilin 1 and 2
RNA	Ribonucleic acid
sAPP	Secreted amyloid protein precursor
SCI	Spinal cord injury
Sox 2	Sex-determining region Y-box 2
STAT3	Signal transducer and activator of transcription 3
SVZ	Sub-ventricular zone
TUNEL assay	Terminal deoxynucleotidyl transferase dUTP nick-end labeling

#### 1 Introduction

Aging is something that all living things must endure; however, how we approach aging as individuals can be vastly disparate due to genetics or environmental conditions. Each time cellular division occurs, telomeres, the protective noncoding deoxyribonucleic acid (DNA) sequences found at the ends of chromosomes on the 3' end, are shortened due to the replication mechanism of the lagging strand. Each cell can undergo a predetermined number of replications before telomeres have been exhausted, at which point the cell receives signaling to become senescent and die. This continual process of telomere shortening and cell death is commonly known as the process of aging.

In the 1980s, an enzyme was discovered that allowed cells to bypass cellular aging and divide without exhausting their telomeres. For the past two decades, researchers have toiled away at unlocking the secrets of this enzyme, telomerase, thought to be the modern-day fountain of youth. Telomerase is a reverse transcriptase containing its own template ribonucleic acid (RNA) molecule, allowing it to add additional telomeres to the ends of chromosomes. Somatic human cells have been found to be severely lacking in telomerase and continue to experience telomere shortening at each division, causing DNA damage and cell death. In contrast, stem cells, cells that are described as any cell that has the ability to renew itself and differentiate into different types of cells, were found to be rich in telomerase.

Due to their high levels of telomerase activity, stem cells are capable of dividing without experiencing telomere shortening. These unique properties allow the body to continue to thrive, as stem cells replace damaged or exhausted somatic cells throughout the body. However, not even stem cells are invincible to the aging process. As the body ages, stem cells become less active, and the rate of cellular turnover decreases. It is the current belief that if the stem cell population could be increased or reactivated, it would be possible to slow the effects of aging and even mitigate many age-related and degenerative diseases.

The limitless possibilities for the use of stem cells have sparked a new era in the field of regenerative medicine and have led the scientific community to recognize stem cell therapeutics as the key to unlocking future treatments for many disabling and currently incurable diseases. Companies and medical institutions around the globe have poured billions into the development of stem cell technologies and are now offering various stem cell-related products and therapies. Many of these technologies are currently in the process of clinical trials to seek approvals from officials, making it possible for treatments to be available in the European Union and United States in the near future.

# 2 Embryonic versus Adult Stem Cell Therapeutics

There are two prominent types of stem cells: embryonic stem (ES) cells, isolated from blastocysts, and adult stem (AS) cells, referring to more specialized stem cells present throughout the body after embryonic development. AS cells have been found in tissues and organs such as brain, skin, heart, liver, gut, testis, blood, and bone marrow. Over the past few decades, it has been widely debated as to which of these sources is the most effective for the development of safe and successful treatments.

ES cells are rapidly self-renewing cells capable of giving rise to any of the three germ layers. Their ability to be kept in culture indefinitely also makes them a valuable research and treatment tool. However, ES cells are only available 4–5 days postfertilization up until the end of embryogenesis (Thomson et al. 1998). The limited time for collection, as well as the availability of willing donors, increases both the difficulty and cost of production. In addition to challenges regarding the availability of this cell population, the use of ES cells has sparked many political debates regarding the ethics pertaining to the use of embryonic tissue, which some view to be the morally adverse destruction of human life. Furthermore, there are two very serious scientific concerns: the bodies' ability to recognize foreign materials – immunorejection can be just as lethal as the potential cure – as well as the presence of tumorigenesis, which occurs when ES cells are transplanted (Wiles and Johansson 1999; Yu et al. 2007).

In contrast, AS cells are more ethically accepted than ES cells since they are isolated from tissues, which does not involve termination of embryogenesis. They are readily available and can be isolated from sources such as umbilical cord, bone marrow, and peripheral blood, thus increasing cost-effectiveness. Also, it allows us to use a patient's own cells to for treatment, known as autologous therapy, which eliminates the concern of immunorejection. Since AS cells are already committed to becoming certain types of cells, there is no issue of tumorigenesis upon transplantation. However, their limited potency also poses a challenge in that easily available AS cells (bone marrow, blood, umbilical cord, etc.) are limited to the treatment of diseases within the scope of the tissue of origin, and cannot be used in instances such as neurological disorders. Nonetheless, many researchers feel that

Embryonic stem cell	ls versus adult stem cells		
Embryonic stem cell	ls	Adult stem cells	
Pro	Con	Pro	Con
Can give rise to all three germ layers	Available only from human blastocysts	Source of cells is more ethically accepted	Limited in its ability to differentiate into cell types
Fast self-renewal	Immunorejection	Easy maintenance	Slow self-renewal
Embryonic stem cells can be kept in culture indefinitely	Ethics issue on destruction of human life	Can be isolated from patients at any time	Limited expansion in culture
	Tumorigenesis	Easy access to the source of cells	
	Limiting funding for R&D	Autologous use of cells to eliminate immunorejection	
	Patent protection on hES cell lines in the USA by the University of the Wisconsin Alumni Research Foundation		

the above concerns regarding the use of ES cells outweigh the benefits and that AS cells may be a much more viable choice for potential therapeutics.

In order to bypass potency limitations of AS cells, researchers have developed induced pluripotent stem (iPS) cells, which are cells that have been reprogrammed to an embryonic-like state. In 2006, researchers used a retroviral vector to transfect mouse fibroblasts with four factors: Oct-3/4, Sox2, Klf4, and with the oncogene Myc. The results from this study showed that these cells represented the morphology of ES cells, as well as expressed ES marker genes, and showed that iPS cells can be directly generated from mouse fibroblasts in culture (Yamanaka 2008), (Meissner et al. 2007; Nakagawa et al. 2008; Okita et al. 2007; Takahashi and Yamanaka 2006). These results were replicated by several independent groups using human somatic cells (Takahashi and Yamanaka 2006; Park et al. 2008) and were also repeated using combinations of Nanog, Oct-4, Sox2, and Lin28 (Yu et al. 2007).

Although these studies represent a major breakthrough in the scientific community for the creation of patient- and disease-specific stem cells, the use of viral vectors and oncogenes in human cells has raised safety concerns. Also, the starting material used for these studies, skin blasts, is highly differentiated with a high level of DNA methylation. This may prevent certain gene expression and stop cells from becoming fully functional. In order to overcome this problem, alternative starting material with less DNA methylation, such as adult stem cells, which have less methylation and may be more flexible, needs to be explored. One such study used mesenchymal stem (MS) cells as the starting material for the induction of pluripotency. Not only was this study successful in creating MS cell-derived iPS cells, but it was done using a single factor, Nanog, as well as a nonviral transfection method (Alvarez et al. 2010). In addition, this study went on to show that the generated iPS cells were capable of developing into cells exhibiting neural cell-like characteristics, demonstrating their ability to transdifferentiate (Alvarez et al. 2010).

These promising studies show incredible potential for the development of treatments for a limitless number of diseases, and with momentum continually building in the field of iPS research, it is not unreasonable to predict that iPS cells will soon be implemented in clinical therapeutic studies.

## 3 Alzheimer's Disease

Alzheimer's disease (AD) is a neurological disorder effecting the central nervous system characterized by the progressive loss of memory and intellectual functioning, typically becoming apparent after age 65 (Cruts and Van Broeckhoven 1998). AD is the most common type of dementia associated with aging, which is now the sixth leading cause of death for those 18 years of age and above and the fifth for those 65 and above. It is expected that 2.6–5.2 million Americans will be diagnosed with AD; if this current trend continues we will have over 16 million Americans diagnosed with AD by the year of 2050. The economic impact is projected to increase from 183 billion for 2011 to 1.1 trillion in 2050 (an executive summary progress report on the CDC Healthy Brain Initiative 2006–2011). As an incurable, progressive, and terminal disease with limited treatment options, this changes the status of what was once considered a rare disorder to a major medical disease that impacts a vast amount of both families and patients (Brookmeyer et al. 2011).

## 4 Alzheimer's Disease Pathology

As AD progresses, cell-to-cell communication is disrupted, ultimately leading to cell death and loss in brain mass. This devastating reduction of cells affects nearly all brain functions. Symptoms of AD may include memory loss which disrupts daily life, challenges in problem solving and planning, difficulty completing common daily tasks, confusion with time or place, difficulty interpreting spatial relationships and visual images, problems with speaking and writing, misplacing objects without the ability to retrace one's steps, poor judgment, withdrawal from social activities or work, and changes in mood and personality.

The mechanism behind AD is still greatly researched and debated. The most widely recognized theory for the pathology of this disease is the amyloid cascade hypothesis, which describes the hallmark formation of plaques. In this model, plaques are caused by the accumulation of beta amyloid ( $\beta$ -amyloid) between neurons.  $\beta$ -amyloid is a normally occurring, sticky protein fragment produced in the brain when it is snipped from amyloid precursor protein (APP) via sequential scission, the  $\beta$ -APP-cleaving enzyme (BACE) (Hussain et al. 1999; Lin et al. 2000;

Sinha et al. 1999; Vassar et al. 1999; Yan et al. 1999; Karran et al. 2011); however, in the AD brain, it is undigested and gathers into hard, insoluble plaques. Another widely studied hallmark of AD is the formation of tangles. Intracellular neurofibrillary tangles (NFT) occur when the tau protein, a highly soluble microtubule-associated protein, which aids in the formation of microtubules (Cleveland et al. 1977; Obulesu et al. 2011), is hyperphosphorylated. Hyperphosphorylated tau is incapable of binding to and stabilizing microtubules, and they therefore collapse, thus creating insoluble twisted fibers inside brain cells (Obulesu et al. 2011; Mandelkow et al. 2007; Larbig et al. 2007; Metcalfe and Figueiredo-Pereira 2010). Some researchers argue that these two hallmarks, plaques and tangles, may occur at the late-stage of the disease, and that they do not address the initial biochemical mechanisms of AD. Other researchers have focused on what drives the loss of connections between brain cells, disruption in cells signaling pathways, and inflammation as alternative possible pathways for AD.

Both risk and deterministic genes have been identified in AD. Four genes have been found to play a crucial role in the onset of Alzheimer's disease: amyloid precursor protein (APP), presenilin-1 (PS-1), presenilin-2 (PS-2), and apolipoprotein E-e4 (APOE4). One crucial pathological condition of AD is the amyloid beta deposition, plaques, in the brain. This occurs through both excess production and miscutting of amyloid beta (A4) precursor protein (APP).

# 5 Current Therapies for Alzheimer's Disease

There is currently no cure for Alzheimer's Disease; however, some strides have been made in slowing down the progression of the disease in the hopes that a viable treatment is on the horizon. Memory loss and confusion, the most debilitating symptoms of AD, are presently being treated using (drugs approved by the FDA) cholinesterase inhibitors (Aricept, Exelon, Razadyne, Cognex, and Namenda). Donepezil (brand name Aricept) is approved for all stages of AD and can present with side effects such as vomiting, nausea, increased bowel movement, and loss of appetite. Galantamine (brand name Razadyne), rivastigmine (brand name Exelon), and tacrine (brand name Cognex) have all been approved for AD presenting as mild to moderate progression. Symptoms for these vary from nausea, vomiting, increased bowel movement, loss of appetite, and possible liver damage. Memantine (brand name Namenda) is for AD presenting as moderate to severe in progression and has side effects such as headache, constipation, confusion, and dizziness. Even though these treatments help in slowing the progression of AD, it tends to equate to putting a Band-Aid on a gashing wound.

#### 6 Stem Cell Replacement Strategies for Alzheimer's Disease

Studies conducted using neural stem (NS) cells isolated from both fetal and adult mammalian cells, (Doetsch et al. 1999; Gould et al. 1999) located within the central

nervous system (Johansson et al. 1999) (CNS), were proliferated in vitro using a variety of culture medium systems (Svendsen et al. 1998) then transplanted in already progressive AD environments with hopes of creating new neuronal connections. These culture systems are crucial in that they contain certain factors that are necessary for successful integration into the host CNS. There are currently established methods of maintaining and expanding NS cells in culture in a serumfree defined medium that contains basic fibroblast growth factor (bFGF) and the epidermal growth factor (EGF) (Brannen and Sugaya 2000), (Fricker et al. 1999). However, there are issues related to NS cells regarding their much shorter telomeres and that they express almost no telomerase after a certain number of passages (Ostenfeld et al. 2000). Other researchers have been successful in keeping these cells in culture long term while maintaining the ability to differentiate into cells that stain positive for beta III tubulin, a neuronal marker, and glial fibrillary acidic protein (GFAP), which is a marker for astrocytes, in neurobasal medium without the addition of exogenous factors. This suggests that these cells, once in the appropriate environment, are able to produce factors on their own. With these studies in mind we are able to expand NS cells in vitro in order to produce a well-defined source of transplantable material to continue investigations into stem cell replacement strategies.

One such study used NS cells that were expanded without exogenous mitogenic factors using serum-free media (Brannen and Sugaya 2000) then injected into the lateral ventricle of both mature (6-month-old) and aged (24-month-old) rats. Cognitive capabilities were tested using a Morris water maze 4 weeks posttransplantation. Results demonstrated a markedly significant increase in cognitive function of memory-impaired rats to that of a level of memory unimpaired or to a level considered normal for the animal's age. Postmortem evaluation by immunohistochemistry showed positive for both beta III tubulin and GFAP. Positively stained neurons were found in the bilateral cingulate, in the parietal cortexes, and in the hippocampus. Morphological identities associate with these locations indicate that the aged brain is still able to guide migration and differentiation patterns. Other researchers have tried to pre-differentiate cells into specific neuronal types in vitro and then transplant to a direct location; however, these efforts showed no host integration or increased cognition (Clarkson 2001). This could be due to the fact that fully differentiated cells are no longer able to migrate and, therefore, cannot successfully integrate into the system. Another issue could be the destruction of tissue produced by injecting cells into a direct location, which could cause an immune response resulting in the destruction of transplanted tissue. These discoveries promote that it is a key element to deliver undifferentiated versus differentiated cells into the brain ventricle instead of using a direct location and overall use of NS cells poses promising opportunities as a stem cell replacement strategy.

Another group used NS cell transplantation in order to reduce cerebral ischemia within the brain. Adult rats were given intracerebral hemorrhages by administering bacterial collagenase intrastriatally. In the study, the NS cell groups showed better function on the rotarod test - a performance test used to evaluate balance and coordination - two weeks posttransplantation and improvement 5 weeks

posttransplantation with the modified limb-placing test assessing sensorimotor integration of the limbs. These results suggest that NS cell transplantation can be a viable treatment format for the restoration of neurological defects (Jeong et al. 2003).

Chemicon, a company that was recently acquired by Millipore, developed two immortalized human neural progenitor cell lines, ReNcell VM and CX. ReNcell VM was derived from the ventral mesencephalon of the fetal brain, whereas CX comes from the cortical region of the fetal brain. Both cell lines share the ability to differentiate into neurons and glial cells along with the ability to continually reproduce themselves. The company further showed that the cells differentiated into high-level neurons and maintained their electrophysiology. With this in mind, these cells have the potential to become a therapeutic treatment for AD (Donato et al. 2007).

Other recent findings promote stroke-based therapies generated through selfrepair of neurons. Induced middle cerebral artery occlusions led to the death of striatal neurons, which then triggered an increase in NS cell production through activation of the dormant NS cells already located in the sub-ventricular zone (SVZ). The newly premature neurons were able to migrate to the location of the lesions and repair it through the production of new neurons (Lindvall et al. 2004). The area of spinal cord injury (SCI) is another topic of interest in the research arena. One such group using isolated human fetal central nervous system (CNS)-derived stem cells grown as neurospheres were transplanted into immunodeficient (scid) mice. The cells were able to survive the transplantation, migrate, and differentiate with minimal GFAP-positive expression. Engraftment into the host was demonstrated through improved locomotor functioning. This provides promising results for hNSCs as a potential treatment for spinal cord injuries (Cummings et al. 2006).

As demonstrated by the abovementioned studies, it is possible the NS cells could provide a possible therapeutic approach to many diseases that currently have no productive or readily available cure. However, these studies are still in their infancy and will need much more elucidation before becoming an acceptable treatment choice and many challenges lay in its way.

# 7 Pharmacological Applications as a Neuroreplacement Strategy

In brain development, production of new nervous system cells occurs in early childhood and fetal stages; however, the adult brain also contains deposits of stem cells all the way into advanced years (Galli et al. 2003). Neurogenesis was first documented in the SVZ (Gould et al. 1999) and the dentate gyrus of the hippocampus (Eriksson et al. 1998; Kuhn et al. 1996; Gage 2000). Specific signaling induces mitogenic factors, such as EGF and bFGF, to direct the division of endogenous neural stem (eNS) cells located in these areas and move them towards differentiation and migration to their final destination. Plasticity of adult eNS cells has also been reported by numerous groups (Eriksson et al. 1998; van Praag et al. 2002).

In the normal brain, the rate of production of proliferating NS cells appears to be balanced by apoptosis, allowing for controlled modeling and maintenance of brain structures. Aging and stress may decrease the proliferation of NS cells (Kuhn et al. 1996), while neuronal production is upregulated in response to brain injury and ischemic diseases (Felling and Levison 2003), suggesting that eNS cells may play a role in repair as well as normal adaptive responses. When the mechanism of selfrenewal and repair is no longer able to cope with the loss of neuronal mass, progression of neurodegenerative diseases occurs, leaving patients to suffer functional brain deterioration and behavior patterns ultimately leading to a decline in the quality of life. The rate of neurogenesis from eNS cells tends to rise or fall in response to physiological conditions, such as aging, diseases, and stress (Kuhn et al. 1996; Felling and Levison 2003; McEwen 2001; Jin et al. 2004), indicating that eNS cell proliferation may be controlled by certain intrinsic factors. Many researchers have isolated factors from the conditioned media of neural stem cells and tested these factors in various combinations. Fibroblast growth factor (FGF) and epidermal growth factor (EGF) have been implicated in neural migration, axon navigation, and synaptogenesis (Guillemot and Zimmer 2011) and have the ability to increase proliferation of neuronal precursor cells (Arsenijevic et al. 2001).

It stands to reason that the development of similar compounds that mimic these conditions would be a valuable tool.

Therefore, these factors have become a major pharmacological target. In one study, researchers purified an autocrine/paracrine cofactor, a glycosylated form of cystatin C (CCg), and combined it with the trophic factor, basic FGF (FGF-2), both in vitro and in vivo. The addition of both of these factors showed stimulated neurogenesis in the dentate gyrus, thus suggesting that neurogenesis may be controlled by both trophic factors and autocrine/paracrine cofactors, namely, CCg (Taupin et al. 2000). Similarly, another study showed that FGF or EGF alone cannot stimulate neurogenesis. In this study, researchers considered the early appearance of insulin-like growth factor (IGF-I) receptors during mouse striatum development and its role in NS cell regulation. Combinations of these three factors were used in order to study the action of IGF-I on striatal NS cell proliferation. The results demonstrated that proliferation was not induced in the presence of neither EGF nor FGF-2 alone; however, the addition of IGF-I resulted in large proliferative clusters of multipotent spheres. This data suggests that IGF-I plays a crucial role in the regulation of NSC activation in conjunction with FGF-2 and EGF (Arsenijevic et al. 2001).

Although these factors seem to play a pivotal role in neurogenesis, there is a major pitfall associated with this treatment course. Since factors are incapable of penetrating the blood-brain barrier (BBB), these types of pharmaceutical would need to be injected directly to the brain. An alternative method of treatment would involve the use of small molecules capable of crossing the BBB, such as antidepressants, shown to increase stem cell proliferation. In one study, human hippocampal cells from untreated subjects with major depressive disorder were used to test the effect of selective serotonin reuptake inhibitors and tricyclic antidepressants on increasing neural progenitor (NP) cells and proliferation in the human dentate gyrus. Results showed that these antidepressants were capable of increasing NP cells in the anterior human dentate gyrus (Boldrini et al. 2009). Another study confirms these findings and reports that chronic treatment with antidepressants increased cell proliferation in the hippocampus of a rodent model (Malberg et al. 2000). Although these finding may be significant for the treatment of depression, the increase of cell proliferation will most likely not be significant enough for the treatment of neurodegenerative diseases such as AD.

One such small molecular compound, a pyrrolopyrimidine (MS-818) (Itoh et al. 1999), may be a promising target for increasing stem cell proliferation. This compound was initially identified for its ability to promote neurite outgrowth in isolated neurons maintained in vitro (Awaya et al. 1993). In another cell culture study using neurons isolated from rodent cortex, MS-818 was reported to be active primarily in the presence of various growth factors, including bFGF, nerve growth factor (NGF), EGF, and insulin-like growth factor (IGF-1) (Sanjo et al. 1998; Torigoe and Awaya 1998; Sugiyama et al. 2002; Watanabe et al. 1998). The mechanism of action of MS-818 is not known, although some evidence implies the activation of the MAPK (mitogen-activated protein kinase) pathway, a cascade that is also activated by peptide growth factors, may play a key role (Sanjo et al. 1998). In the same report, it was suggested that MS-818 promoted the survival of rodent cortical neurons by reducing the rate of apoptosis, as measured by TUNEL assays. MS-818 has been tested in isolated animal models, axon growth in mice (Torigoe and Awaya 1998), and muscle regeneration in rats (Sugiyama et al. 2002), with evidence of but no mechanistic details relating to function.

Recent studies have shown that MS-818 induces the proliferation of endogenous NSCs, which may explain mechanism of this compound's function (Sugaya 2005; Sugaya and Merchant 2008). In these studies, they have shown that MS-818 increased NS cell population 600 % in the cortex of aged rodent models. It was also found that MS-818 treatment improved behavior of Parkinson's disease model in rotarod test with association of increased proliferation of NS cells. Although toxicity studies using MS-818 have not been reported, the compound was not found to have obvious side effects in rodents in these studies, and even at the high dose, it may be a good candidate for a pharmacological treatment of AD (Sugaya et al. 2007, 2006; Kwak et al. 2006a).

# 8 Obstacles for Neuroregeneration Therapy for Alzheimer's Disease

The ability to increase eNS cells populations via a small molecular compound, MS-818, as well as transplanted NS cells in order to recover cognitive function in aged rats has been demonstrated. However, we may need to consider that AD pathologies may affect NS cells and reduce the efficacy of these neuroregenerative treatments. Over the past decade, scientists have uncovered much about AD pathologies such as the APP and tau defects, along with other related gene mutations, but how these modify the AD environment needs to be elucidated in order to the successful stem cell therapy of this disease.

One such example of the challenges associated with the diseased environment of the AD brain is induction of glial differentiation of NS cells as a result of high concentration of APP in the environment. In a recent study, NS cells were transplanted in the brain of APP transgenic mice where human APP is highly expressed to find whether NS cell can be used to treat AD. The hope of this study was that the transplanted cells would differentiate into functional neurons. However, this research lead to the conclusion that the high concentration of APP induced glial differentiation of NS cells instead of the much needed neuronal differentiation (Marutle et al. 2007; Kwak et al. 2011, 2010, 2006b, 2006c; Keilani and Sugaya 2008). It has shown that recombinant secreted amyloid precursor protein (sAPP) dose dependently differentiated NS cells into astrocytes.

A further study was conducted in order to elucidate the pathway of APP induced glial differentiation through the activation of interleukin-6 (IL-6) and glycoprotein (gp130) signaling pathways, which are associated with inflammation. Gliogenesis, the formation of glial populations derived from multipotent neural stem cells, is modulated through certain external factors: leukemia inhibitory factor (LIF), ciliary neurotrophic factor (CNTF), bone morphogenetic protein (BMPs), and Notch along with integral factors neurogenin 1 (Ngn1), neurogenin (Ngn2), and deoxyribonucleic acid (DNA) methylation. It has been revealed that all of these above factors are responsible for glial differentiation through the activation of the IL-6/gp 130 signaling pathway. Ligands, LIF and CNTF, bind to receptors and subsequently phosphorylates gp130, Janus kinase (JAK), and signal transducer and activator of transcription 3 (STAT3) pathways. STAT3 then translocates into the nucleus in order to stimulate gene expression of the target gene such as glial fibrillary acidic protein (GFAP). The IL-6/gp130 signaling pathway is modulated by an autoregulatory loop which eventually coordinates with other pathways related to glial differentiation, such as Notch and BMP {Kwak, 2010 #1990}. This helps to confirm previous studies showing that sAPPa in high concentrations works as a gliogenic factor in stem cell differentiation.

The APP gene is located on chromosome 21. Trisomy of chromosome 21 is responsible for most cases of Down syndrome (DS). DS patients have shown AD symptoms as early as 40 years of age (Schnabel 2011); this may be due to the extra copy of chromosome 21, which produces an extra copy of APP gene also. Svendsen (Svendsen et al. 2001) found that DS patient's do not produce normal level of neurons and even those that are produced, the morphology is altered (Sugaya et al. 2007; Schnabel 2011; Svendsen et al. 2001; Svendsen and Caldwell 2000). When APP genes are overexpressed in the normal NS cells, they show exactly the same, less production of neurons and altered morphology.

One of the symptoms in aging and AD is the loss of AS cells, for it is this cell population that will ultimately be needed to replace neuronal functioning within the AD brain. In a healthy individual, AS cells will remain quiescent within the body in a nondividing state until they are called upon to perform duties for the purpose of tissue maintenance, disease-related repairs, or when injury has occurred. The slow exhaustion of these cells over an extended time course may lead to the natural symptomatic aging process; however, under diseased conditions such as AD, the number of AS cell may decline more rapidly than normal aging. Abovementioned APP functions to induce glial differentiation of NS cells and induce premature differentiation of NS cell population in the AD or DC. Thus, regulation of APP level in the AD may be a useful prevention therapy as well.

# 9 (+)-Phenserine's Ability to Overcome Pathological Effect of Alzheimer's Disease

Before treatment for AD can begin, modification must be made to the environment which will allow for the differentiation of neuronal cells. The challenge is in discovering how to go about this without disrupting other normal functioning. There are several cholinesterase inhibitors (ChEIs) currently available for the treatment of mild to moderate AD (e.g., Donepezil, Galantamine, and rivastigmine). These ChEIs have been found to improve memory, thinking, language, and judgment through the increase of acetylcholine levels in the brain. It is believed that this increase of acetylcholine may increase cognition by compensating for loss of functioning brain cells as stated by an Alzheimer's Association report regarding FDA-approved treatments for Alzheimer's disease. Additionally, ChEIs have been found to regulate APP levels. (-)-Phenserine is one such inhibitor known to decrease levels of APP. However, the dosing of (-)-phenserine has presented a challenge. In order to suppress the appropriate amount of APP and A $\beta$  levels, the concentration of (-)-phenserine must be significantly increased. At such high concentrations, (-)-phenserine has been known to produce severe side effects, such as increased production of acetylcholine, resulting in the autonomic nervous system to become unbalanced. In order to efficiently regulate APP levels and bypass the side effects associated with (-)-phenserine, an isoform, (+)-phenserine, was developed. When using this isoform, it is possible to give the drug at the required doses without eliciting the associated side effects (Marutle et al. 2007).

## 10 Conclusion

Aging is the highest risk factor for AD. Although the role of stem cells in the adult remains unclear, reduction of the stem cell population associated with aging may significantly affect brain function. If the other factors are combined, such as a high concentration of APP or inflammation in the brain, the reduction of NS cells during the aging could be accelerated. Thus, augmentation of the stem cell population or transplantation of stem cells may help to find a cure for AD. However, we have to always consider the effect of the pathological condition of AD on stem cell biology as the mature neurons may not be able to incorporate into the host brain as functional cells. We have iPS cell technology to produce neural cells from somatic cells and small molecular compound which increases endogenous stem cell population, which allow us to perform autologous regenerative therapies. As our knowledge increases in clear understanding the pathological changes in AD brain, we will be able to fight against aging and AD.

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# Tardive Dyskinesia: Outcome of Antipsychotic Treatment and Brain Damage?

# Richard M. Kostrzewa, John P. Kostrzewa, and Ryszard Brus

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R.M. Kostrzewa (🖂)

Department of Biomedical Sciences, Quillen College of Medicine, East Tennessee State University, Johnson City, TN, USA e-mail: KOSTRZEW@mail.etsu.edu

J.P. Kostrzewa

Department of Biomedical Sciences, Quillen College of Medicine, East Tennessee State University, Johnson City, TN, USA

North Alabama ENT Associates, Huntsville, AL, USA e-mail: jpkostrzewa@hotmail.com

R. Brus

Department of Nursing, High School of Strategic Planning, Dabrowa Górnicza, Poland e-mail: hrbrus@wp.pl

#### Abstract

Tardive dyskinesia (TD), marked by abnormal involuntary movements and frequently expressed as perioral activity, represents an adverse outcome of prolonged antipsychotic therapy, occurring in approximately 5 % of patients per treatment year. Although neuronal mechanisms underlying TD are largely unknown, more recent experimental studies in animal models of TD are providing insight into the neuronal mechanisms associated with TD and implicating newer treatment approaches. It is now evident that a predominance in the ratio of dopamine (DA)  $D_1:D_2$  receptor (R) activation accounts for induction of perioral movements in rodent models of TD, in nonhuman primate models of TD, and in humans with TD. Experimentally, TD is produced in animal models of TD, in a manner analogous to that by which TD is produced in humans – by continuous and prolonged administration of a DA D<sub>2</sub>R antagonist (i.e., an antipsychotic drug). More recently, in a rodent model of TD, it has been shown that a lesion of dopaminergic - mainly nigroneostriatal neurons reduces the time latency for occurrence of TD, also increases the severity of perioral activity, and results in permanence of TD after complete removal of D<sub>2</sub>R antagonist treatment. The induction of perioral activity is related to DAR supersensitivity but unrelated to numbers of  $D_1R$  and  $D_2R$  in the neostriatum, a brain region associated with perioral activity. More apropos, serotoninergic systems are now recognized as having a greater role in effecting perioral activity, and it appears that 5-HT<sub>2C</sub> receptor antagonists are most effective in abating perioral activity in a rodent model of TD. These processes and mechanisms, topics addressed in this chapter, highlight a newer understanding of mechanisms underlying TD and provide insight into new approaches towards treatment of TD in humans.

#### Keywords

 $5-HT_2$  receptor • 5-Hydroxytryptamine • 5,7-Dihydroxytryptamine • 6-Hydroxydopamine •  $D_1$  receptor •  $D_2$  receptor • Dopamine • Haloperidol • Perioral movements • Receptor supersensitivity • Serotonin • Tardive dyskinesia • Vacuous chewing

#### 1 Introduction

On the website of the National Institute of Neurological Disorders and Stroke (NINDS) (http://www.ninds.nih.gov/disorders/tardive/tardive.htm), tardive dyskinesia (TD) is classified as an extrapyramidal syndrome arising from long-term use of neuroleptic agents (i.e., antipsychotics). TD is characterized by repetitive involuntary purposeless movements of specific muscle groups which may include fingers, arms, legs, and trunk but generally presents as oral dyskinesia (typically, chewing movements) which may include lip smacking and tongue protrusion. A person with TD may appear to be chewing gum.

TD arises in humans, nonhuman primates (Blanchet et al. 2012), and rodents by chronic (exclusive) treatment with a dopamine (DA)  $D_2$  receptor (R) antagonist (Saifee and Edwards 2011) – the class of agents oft used for treating schizophrenia. The incidence rate of TD is approximately 5 % per year of treatment with a "classical" (first-generation) antipsychotic drug (e.g., haloperidol) (Kane et al. 1985) and two to threefold less so when an atypical antipsychotic agent (e.g., clozapine) is used (Tarsy et al. 2011). TD tends to be long lived, sometimes lifelong, and is usually refractory to treatment. TD remains a serious problem today. Tetrabenazine is the only approved drug for TD, but a variety of other drugs is often attempted.

#### 2 Multineuronal Associations with Tardive Dyskinesia

Although TD arises from prolonged administration of a  $D_2R$  antagonist, it is recognized that many different neuronal phenotypes may have a role in the development or persistence of TD (Casey 1987; Egan et al. 1995; Jeste and Caligiuri 1993; Gong et al. 1992; Gunne and Häggström 1983; Knable et al. 1994; Kostrzewa 1995; Waddington 1990). Cholinergic (Kostrzewa and Neely 1993; Rupniak et al. 1983; 1985; Salamone et al. 1990), GABA-ergic (Gunne et al. 1988; Lloyd et al. 1985; Mithani et al. 1987; Tamminga et al. 1979), and serotoninergic (Gong and Kostrzewa 1992; Gong et al. 1992) involvements have been implicated in TD. Nevertheless, the focus in this chapter will remain on dopaminergic systems and the recently uncovered important role of serotoninergic systems in TD.

## 3 Oral Dyskinesia Arising in Rodents from Acute Dopamine Receptor Agonist or Antagonist Treatments

Acute administration of a DA  $D_2R$  antagonist to rats or mice results in the onset of oral dyskinesias, which are prominent over the following hour or more. The short-lived effect is related to the dose of the  $D_2R$  antagonist. Similarly, acute administration of a DA  $D_1R$  agonist is associated with a short-lived and dose-related induction of oral dyskinesia. These correlated findings give credence to the possibility that oral dyskinesia may be related to an imbalance in  $D_1R/D_2R$  activation by the endogenous neurotransmitter, DA (Rosengarten et al. 1983). Moreover, when a  $D_2R$  antagonist is administered chronically, as per animal modeling of TD, or when psychiatric patients are treated long term with an antipsychotic drug, one must consider the long-term effect of such treatment on the overall balance of  $D_1R/D_2R$  activation – and the effect this might have on the origin of TD.

## 3.1 Experimental Induction of Oral Dyskinesia in Rodents by Acute Dopamine D<sub>1</sub>R Agonist Treatment or D<sub>2</sub>R Antagonist Treatment

Antipsychotics and  $D_2R$  antagonists in general induce the occurrence of spontaneous oral activity – the so-called oral dyskinesias in humans and the so-called vacuous chewing movements in rodents (Clow et al. 1979; Iversen et al. 1980; Waddington and Gamble 1980). In contrast, DA  $D_1R$  agonists induce oral activity (Arnt et al. 1987; Koshikawa et al. 1987; Levin et al. 1989; Molloy and Waddington 1988; Murray and Waddington 1989; Rosengarten et al. 1983; 1986a, b; Rupniak et al. 1985), and it is the presumed balance in  $D_1R/D_2R$  activation that is considered to be important for this behavioral response (Rosengarten et al. 1983). For example, in rodents, there is an increase in oral activity when there is an increase in  $D_1R/D_2R$ activation, as occurs with  $D_2R$  antagonists, also after  $D_1R$  agonist treatment (Johansson et al. 1987; Molloy and Waddington 1987; 1988; Rosengarten et al. 1983), and in rat strains with reduced numbers of  $D_2R$  (Rosengarten et al. 1986a, b).

## 3.2 Effect of Neonatal Dopaminergic Denervation on Induction of Oral Dyskinesia in Rodents by Dopamine D<sub>1</sub>R Agonists or D<sub>2</sub>R Antagonists

When rats are lesioned perinatally with the neurotoxin 6-hydroxydopamine (6-OHDA; desipramine pretreatment), repeated  $D_1R$  agonist treatments "prime" DA  $D_1R$ . Initial  $D_1R$  agonist treatments have no marked effect, but repeated  $D_1R$  agonist treatments permanently sensitize  $D_1R$ , so that exaggerated behavioral responses are observed. This process is known as homologous priming of  $D_1R$ , and the resulting  $D_1R$  supersensitization is lifelong (Breese et al. (1985, 1987). Also, repeated  $D_2R$  agonist treatments of 6-OHDA-lesioned rats can induce the  $D_1R$  supersensitization, a process known as heterologous  $D_1R$  priming (Criswell et al. 1989).

When rats so-lesioned neonatally with 6-OHDA were challenged with a single dose of a  $D_2R$  antagonist, there was an increased in perioral movements when compared to the effect in non-lesioned control rats (Kostrzewa and Hamdi 1991). In an analogous manner the first challenge dose of a  $D_1R$  agonist to 6-OHDA-lesioned rats in adulthood surprisingly produced an increased number of perioral movements in comparison to non-lesioned control rats (Kostrzewa and Gong 1991). Thus, in contrast to the latent  $D_1R$  supersensitivity for locomotory and stereotyped activities that required unmasking by repeated  $D_1R$  agonist treatments (Breese et al. 1985; 1987), there was overt supersensitization of  $D_1R$ for  $D_1R$  agonist induction of perioral activity. Moreover, in the 6-OHDA-lesioned rats, perioral activity was induced by a  $D_1R$  agonist dose that was 100–1,000 times lower than was needed for induction of perioral activity for this effect developed if the nigroneostriatal dopaminergic fiber lesioning by 6-OHDA occurred within the first week postbirth (Kostrzewa et al. 1993) and if neostriatal DA depletion was 98.5 % or more (Gong et al. 1993). This  $D_1R$  supersensitization was also lifelong. Moreover, the effect was unaccompanied by an increase in the  $B_{max}$  for  $D_1R$  (i.e., relative number of  $D_1R$ ) or change in the  $K_d$  (i.e., relative affinity of the receptor for an agonist) (Gong et al. 1994).

Perinatal 6-OHDA treatment which produces near total destruction of nigroneostriatal dopaminergic neurons (Doucet et al. 1986) leaves the ventral tegmental nucleus largely intact and able to provide dopaminergic innervation to limbic regions (Snyder et al. 1986; Fernandes Xavier et al. 1994). Thus, the important element of the above studies is that a lesion of dopaminergic innervation to the neostriatum resulted in overt  $D_1R$  supersensitization for the induction of oral activity (Kostrzewa 1995; Kostrzewa et al. 1998, 2003, 2008).

## 4 Oral Dyskinesia Arising in Rodents from Acute Serotonin Agonist Treatment

When neonatally 6-OHDA-lesioned rats were challenged in adulthood with the serotonin (5-HT) 5-HT<sub>2</sub>R agonist m-chlorophenylpiperazine (mCPP), the initial mCPP dose produced an enhanced increase in the number of perioral movements in comparison to that of intact control rats (Gong and Kostrzewa 1992). This finding and others indicate that  $5-HT_2R$  is supersensitized by 6-OHDA lesioning (El Mansari et al. 1994). The effect was not replicated by 5-HT<sub>1A</sub>R agonist 8-OH-DPAT  $[(\pm)$ 8-hydroxydipropylaminotetralin] or 5-HT<sub>1B</sub>R (7-trifluoromethyl-4(4-methyl-1-piperazinyl)-pyrrolo agonist CGS-12066B [1,2-alquinoxaline]); nor was the effect attenuated by 5-HT<sub>1A/1B</sub>R antagonist pindolol, nor by the predominate 5-HT<sub>2A</sub>R antagonist ketanserin, nor by 5-HT<sub>3</sub>R antagonist MDL 72222 (3-tropanyl-3,4-dichlorobenzoate) (Gong et al. 1993). However, the selective 5-HT<sub>2</sub>R antagonist mianserin blocked the effect of mCPP, indicating that 5-HT<sub>2</sub>R, perhaps 5-HT<sub>2</sub>CR specifically, represents the subtype that is overtly supersensitized by 6-OHDA lesioning and associated with perioral movements (Gong et al. 1993).

In fact, the number of perioral movements after mCPP treatment was greater than that following  $D_1R$  agonist treatment (Gong et al. 1992). Mianserin pretreatment, in addition to blocking the effect of the 5-HT<sub>2</sub>R agonist mCPP, blocked the effect of a  $D_1R$  agonist on perioral activity (Gong et al. 1992), and the dorsal striatum was one site for this interaction (Plech et al. 1995; Salamone et al. 1990). By contrast, a  $D_1R$  antagonist did not block the effect of mCPP (Gong et al. 1992). These findings indicate that DAR supersensitivity for perioral activity is mediated via a serotoninergic system and notably at the 5-HT<sub>2</sub>R (Gong et al. 1993).

5-HT fiber hyperinnervation of neostriatum and the forebrain is prominent in rats lesioned perinatally with 6-OHDA (Berger et al. 1985; Snyder et al. 1986; Mrini et al. 1995; see Kostrzewa and Descarries 1998), and this is accompanied by an approximate 30–60 % increase in the number of 5-HT<sub>1A</sub>R, 5-HT<sub>1B</sub>R, 5-HT<sub>1nonAB</sub>R, 5-HT<sub>2A</sub>R, and 5-HT<sub>2C</sub>R (Radja et al. 1993). Therefore, to further explore the effect of 5-HT innervation on perioral activity, the neurotoxin 5,7-dihydroxytryptamine

(5,7-DHT) was coadministered with 6-OHDA to perinatal rats which were then studied as adults. In rats co-lesioned with 6-OHDA and 5,7-DHT, the supersensitized perioral response to a  $D_1R$  agonist was eliminated, indicating that 5-HT fibers indeed mediate the  $D_1R$  agonist effect on perioral movements (Brus et al. 1994).

## 5 Abatement of Oral Dyskinesia in a Rodent Model of Tardive Dyskinesia

Although acute effects of  $D_1R$  agonists on perioral activity are attenuated by acute D<sub>1</sub>R antagonist treatment, a 5-HT<sub>2</sub>R antagonist is just as effective (Gong et al. 1994). In the rodent model of TD, produced by chronic administration of a D<sub>2</sub>R antagonist – over a period of months – acute treatment during the haloperidol-withdrawal phase with a D<sub>1</sub>R antagonist does not reduce the incidence of perioral movements (Kostrzewa et al. 2007). Nor does a D<sub>2</sub>R antagonist (spiperone), muscarinic receptor antagonist (scopolamine), histamine H<sub>1</sub>R antagonist (cyproheptadine), opioid mu receptor agonist (morphine) or antagonist (naloxone), adenosine A<sub>2A</sub>R antagonist (theophylline), alpha-adrenoceptor antagonist (phentolamine, phenoxybenzamine), beta-adrenoceptor antagonist (propranolol); neither a 5-HT<sub>1A</sub>R agonist (pindolol) nor 5-HT<sub>2A</sub>R antagonist (ketanserin). However, agents with 5-HT<sub>2C</sub>R antagonist activity were all effective in acutely reducing the number of perioral movements in the rodent model of TD, produced by vear-long administration of a  $D_2R$  antagonist. Thus, mianserin, mesulergine, ritanserin, and clozapine each acutely reduced spontaneous oral activity during 8 months of the haloperidol-withdrawn phase in the rodent model of TD (Kostrzewa et al. 2007).

Despite the fact that neural reorganization that occurs consequent to monthslong continuous administration of a  $D_2R$  antagonist, it appears that it is an action mediated by serotoninergic 5-HT<sub>2</sub>R that is most consequential (Gong et al. 1992). This is somewhat surprising, because 5-HT innervation in the brain is considered to be an afferent input, not an efferent output system. 5-HT innervation in the brain arises from the brainstem in the dorsal and medial raphe nuclei (see Kostrzewa et al. 1998). Also, from studies in rodents it is recognized that perinatal 6-OHDA destruction of dopaminergic nerves results in supersensitization of DA D<sub>1</sub>R as well as 5-HT<sub>2</sub>R (Gong et al. 1992; Brus et al. 1994). Therefore, in TD it may be that supersensitized 5-HT<sub>2</sub>R has the major influence on the induction of spontaneous perioral activity.

## 6 Tardive Dyskinesia Arising in Rodents from Chronic Dopamine Receptor Antagonist Treatment

Waddington and colleagues (Waddington and Gamble, 1980; Waddington et al., 1983) produced the first rodent model of TD by administering haloperidol in

drinking water to rats for 6 consecutive months. An increase in spontaneous perioral movements developed only at 4 months, thus fulfilling the characteristic "tardive" element of TD (i.e., tardive signifies an onset of symptoms after chronic drug treatment). The increase in spontaneous oral activity persisted for as long as haloperidol continued to be administered, then waned and reverted to normal over a period of a month (see also Waddington 1990). Ultimately, Ellison and See (1989) showed that the form and periodicity of oral dyskinesias in haloperidol-treated rats was similar in form to those of humans with TD. A disadvantage to this model of TD was that the high level of oral activity in rats dissipated over a period of 1 month, back to control level, after haloperidol was withdrawn. This was unlike the persistence of perioral activity in people with TD.

Gunne et al. (1982) and Glassman and Glassman (1980) had observed that there is an increase in neuroleptic-evoked perioral activity in rats following damage to the brain. In the interest of developing a somewhat more novel model and possibly permanent model of TD, it occurred that DA denervation, as occurs with 6-OHDA lesioning, may be an important element in both the prominence and persistence of oral dyskinesia in rodents. Accordingly, we perinatally lesioned rats with 6-OHDA, then added haloperidol to drinking water when these rats were 2 months of age. In these largely DA-denervated rats, an increase in spontaneous oral activity occurred at 3 months, one full month earlier than in intact rats receiving haloperidol via drinking water. Also, the number of perioral movements induced in 6-OHDAlesioned rats by chronic haloperidol was twice the level for chronic haloperidolexposed intact rats. At 1 year the neostriatal Bmax for the D<sub>2</sub>R antagonist  $[^{3}H]$ raclopride was increased  $\sim 25$  % in 6-OHDA-lesioned rats and  $\sim 65$  % in haloperidol-treated rats (both intact and 6-OHDA-lesioned). Notably, when 6-OHDAlesioned rats were withdrawn from haloperidol at 1 year, increased perioral activity did not revert to control levels. The high level of perioral activity observed during the haloperidol phase continued to persist in the group during the withdrawal phase until the study was terminated 8 months later. Further, it is notable that there was no increase in the neostriatal  $D_2R$  (i.e.,  $B_{max}$  for [<sup>3</sup>H]raclopride binding to striatal homogenates) during the haloperidol-withdrawal phase in which there was a persistent high level of perioral activity (Huang et al. 1997). Andersson et al. (1990) have shown that humans with TD do not have an increase in  $D_2R$  in the brain.

## 7 Neural Mechanisms Attending Animal Modeling of Tardive Dyskinesia: Relevance to Human Tardive Dyskinesia

The multiple animal modeling studies relating to TD indicate the following neural mechanism relating to TD. The balance of DA  $D_1R/D_2R$  activation appears to be of critical importance. Both  $D_1R$  agonists and  $D_2R$  antagonists promote oral activity. However, acute effects do not represent a modeling of TD.

TD is, by definition, a "tardive" effect, one that develops over a period of months during prolonged or chronic treatments, specifically with a  $D_2R$  antagonist. TD appears to be induced by prolonged imbalance of  $D_1R/D_2R$  activation, consequent to a tip in the balance towards  $D_1R$  overactivation, possibly in neostriatum or other regions of the forebrain. The nigroneostriatal dopaminergic tract appears to relate most closely to TD. A remodeling of neural circuitry in the brain, over a period of months, represents an adaptive mechanism to the imbalance of  $D_1R/D_2R$  activation, which ultimately results in induction of spontaneous perioral movements. The neural adjustments and reorganization involved in the process are still a mystery.

Nevertheless, prolonged overactivation of DA  $D_1R$  appears to be the genesis for TD. Furthermore, it is apparent that damage to brain circuitry can accelerate the appearance of TD and impact on its permanence. From the most recent experimental studies, it is obvious that damage to dopaminergic innervation to neostriatum promotes earlier onset of TD and permanence. From studies by Gunne et al. (1988), it is possible that damage to other neural circuits in the brain can have similar influence on TD.

## 8 Implications for Treatment of Tardive Dyskinesia in Humans

Because chronic  $D_2R$  antagonist treatment of both humans and rodents produces TD, it is reasonable to assume that similar neural adaptive mechanisms occur in the brain of both rats and humans. Also, because nonspecific injury in the brain (see Gunne and Häggström 1983) as well as nigrostriatal dopaminergic damage confer permanence to the TD produced by prolonged  $D_2R$  antagonist treatment, it is possible that in those patients who develop lifelong TD, there is residual brain damage or dopaminergic fiber degeneration that impacts of the  $D_2R$  antagonist treatment. It is speculated that even a mild brain injury could have occurred earlier in life (i.e., head trauma) or perhaps dopaminergic innervation of the brain declines with aging. The latter event is thought to occur and thereby account for the origin of Parkinson's disease in aged humans (Kish et al. 1992).

Moreover, an increase in the number of neostriatal DA  $D_2R$  is not correlated with high levels of perioral movements, as noted by Huang et al. (1997). Studies by Breese et al. (1985) and Gong et al. (1994) established that DA  $D_1R$  supersensitivity is not necessarily accompanied by an increase in the number of  $D_1$  receptors. Also, DA  $D_2R$  supersensitivity is not correlated with an increase in neostriatal DA  $D_2R$  number in rodents (Kostrzewa and Brus 1991) nor in humans with TD (Andersson et al. 1990). Therefore, the process of receptor supersensitization is not reliant on changes in DA receptor number or affinity. Regardless of underlying factors for development of TD, the animal experimental findings implicate 5-HT<sub>2</sub>R antagonists as reasonable agents for the treatment of TD (Kostrzewa et al. 2007).

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# **Pathogenesis of Alzheimer's Disease**

# Rudy J. Castellani and George Perry

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#### Abstract

Alzheimer disease (AD) is a clinically progressive decline in cortical function involving memory and executive function and pathologically defined by two hallmark lesions, the senile plaque and the neurofibrillary tangle. Nearly 30 years ago, these hallmark lesions were purified and their protein constituents identified by quantitative analysis, which led in turn to a substantial expansion of knowledge, as well as optimism about the ultimate success of targeted therapy. Unfortunately, despite copious facts of new knowledge of the biochemical cascades that produce these protein abnormalities, there has been no meaningful progress toward disease modification for AD, a disease only found in humans. The repeated failures in this regard have been attributed to tardiness in intervention rather than instead an overdue need for a paradigm shift. The ruling theories for AD pathogenesis have their root in lesion removal. We argue that the lack of progress may instead reflect the evolving concept that pathological

R.J. Castellani (🖂)

G. Perry

Division of Neuropathology, University of Maryland, Baltimore, MD, USA e-mail: rcastellani@som.umaryland.edu

Department of Biology and UTSA Neurosciences Institute, The University of Texas at San Antonio, San Antonia, TX, USA e-mail: George.Perry@utsa.edu

lesions, be they plaques, tangles, or soluble low-n protein species, signify mechanisms of neuroprotection, in response to a decade-long adaptation to an aging-hostile environment. The lesions themselves may similarly be a manifestation of neuroprotection, and likewise the targeting of such lesions as the offending agents is done at the very real risk of disrupting homeostasis and the body's attempt at fighting disease. Rather than simply shifting the same ideas and interventions to an earlier age or stage of disease, a broadening of the scope of treatment efforts to working with rather than against the biological processes of the brain, and the realities of repeated negative data, should be accepted both in theory and in practice.

#### Keywords

Aβ • Alzheimer disease • Amyloid-beta • Tau

#### 1 Introduction

Alzheimer disease (AD) is complex and heterogeneous, differing in many respects from case to case, including age at onset, clinical signs and symptoms, presence or absence of various and numerous risk factors, extent and distribution of neuropathology, and genetic alterations including germline mutation and susceptibility alleles. Numerous hypotheses and the treatment implications inherent in them are not unexpected, although the success in treatment has, to date, been absent. Whether or not this failure reflects the complexity of the disease, the now numerous failures in the face of almost limitless knowledge of biochemical facts suggest that prevailing paradigms are significantly, if not fatally, flawed and not salvageable by overlooking theoretical flaws (Castellani et al. 2008a, b, 2009; Castellani and Smith 2011).

Presently, standard thought equates protein constituents of insoluble pathological lesions or their assembly intermediates with toxicity (Hardy and Higgins 1992). AD brains, for example, accumulate neurofibrillary tangles (the major discovery made by Alzheimer in 1906), which are comprised of phosphorylated tau disposed in insoluble fibrils, and those fibrils are derived from soluble tau species which are toxic to neurons in vitro). Thus, the protein "cascade" that produces phosphorylated tau is deleterious (Spires-Jones et al. 2009). Alternatively, one could say that AD brains accumulate amyloid plaques, which contain amyloid beta-protein (A $\beta$ P) in insoluble fibrils of "amyloid"; the amyloid is itself derived from soluble toxic intermediates of A $\beta$ , which are toxic to neurons. The cascade that produces these soluble intermediates is therefore inherently deleterious (Selkoe 2008). In addition, often included in discussions of toxic soluble intermediates is a direct attack on the synapse, closing the loop so to speak on the process in that lesions are tied directly to cognition, although the degree of separation from the in vivo environment is unclear (Tanzi 2005). To be sure, much work needs to be done (Benilova et al. 2012).

#### 2 Amyloidosis in the Brain

A $\beta$  was initially identified through purification of insoluble pathological lesions – senile plaque cores and cerebral vasculature involved by amyloid angiopathy (Glenner and Wong 1984; Masters et al. 1985). Subsequent identification of A $\beta$  as a metabolic product of amyloid  $\beta$ -protein precursor (A $\beta$ PP), its localization to the long arm (q) of chromosome 21, the identification of familial AD kindreds with pathogenic A $\beta$ PP mutations, and the increased AD pathology in Down's syndrome patients who carry an extra copy of chromosome 21 produced a compelling argument in favor of the so-called amyloid cascade as the pathogenic mediator of disease (Castellani et al. 2008a). The cascade was further substantiated by the characterization of presenilin proteins, part of the notch signaling complex with  $\gamma$ -secretase activity (see below) and therefore inherently amyloidogenic.

A $\beta$  is derived from A $\beta$ PP, an integral membrane protein on chromosome 21q21 (Vetrivel and Thinakaran 2006; Wilquet and De Strooper 2004; Ling et al. 2003). Full expression of A $\beta$ PP produces a cytoplasmic tail, a transmembrane domain, and a large extracellular domain, although only a fraction of newly synthesized A $\beta$ PP molecules reach the cell surface. A $\beta$ PP further exists as multiple alternatively spliced isoforms, three of which predominate: two isoforms A $\beta$ PP770/751 contain the Kunitz protease inhibitor domain (exon 7) within the extracellular portion and are the predominant forms in cells other than neurons; A $\beta$ PP695 is devoid of the Kunitz protease inhibitor domain and is the predominant form in neurons.

Cleavage of A $\beta$ PP by either  $\alpha$ -secretase or  $\beta$ -secretase produces soluble N-terminal fragments A $\beta$ PP, and C83 and C99 membrane-bound C-terminal fragments, respectively. Further cleavage by  $\gamma$ -secretase leads to the release and secretion of nonpathogenic p3 peptide (previous  $\alpha$ -secretase cleavage) and A $\beta$  (previous  $\beta$ -secretase cleavage). Moreover, depending on the precise site of  $\gamma$ -secretase cleavage, different lengths of A $\beta$  are produced, varying from 38 to 43 amino acids. The 42-amino acid form, A $\beta$ 42, has a greater tendency to form fibrils in vitro compared to other forms, has a tendency to deposit in brain parenchyma in the form of plaques relative to other species, and has traditionally been regarded as the "pathogenic" A $\beta$  species. A $\beta$ 40 is generally considered benign as well as the principle component of A $\beta$  that is deposited in cerebral vessels in cerebral amyloid angiopathy, particularly cerebral amyloid angiopathy type 2, which tends to spare cortical capillaries. A $\beta$ 42 synthesis and deposition represents the basis for the amyloid cascade hypothesis and therefore the theoretical underpinning of most clinical trials attempted to date.

The constituents and cell biology of  $\gamma$ -secretase have proven a challenge, although these questions were answered in part with the appearance of familial early-onset AD linked not to A $\beta$ PP mutations but to so-called presenilins 1 and 2, presenilin 1 mutations being much more common. Presenilin mutations markedly increase A $\beta$  deposits as well as phospho-tau accumulation in affected patients (Citron et al. 1997; Duff et al. 1996; Scheuner et al. 1996) and have been linked to  $\gamma$ -secretase activity (Maiorini et al. 2002). On the other hand, despite the presumption of presenilin as an important component of the multimeric secretase

complex, the biochemical mechanism of presenilin action is largely unknown. During development, presenilins appear to cleave a transmembrane protein termed notch, which in turn is a transcriptional activator of gene involvement in cellular differentiation (De Strooper et al. 1999). PS1 and PS2 have also been found to be involved in a range of biological processes, including cell adhesion, G-protein-mediated signal transduction, and unfolded protein response (Baki et al. 2001; Smine et al. 1998; Niwa et al. 1999) (Baki et al. 2001; Smine et al. 1998; Katayama et al. 1999; Niwa et al. 1999). Nicastrin has also been shown to interact strongly with the presenilins and appears to be required for normal notch signaling in *Caenorhabditis elegans* (Yu et al. 2000).

APP cleavage with generation of A $\beta$  fragments also differs as a function of cellular subcompartment. At the cell surface, A $\beta$ PP is proteolytically processed, primarily by  $\alpha$ -secretases, resulting in shedding of the majority of the extracellular domain. Rapid and efficient internalization is mediated by a "YENPTY" internalization motif (Vetrivel and Thinakaran 2006). Once endocytosed, A $\beta$ PP may be recycled to the cell surface, degraded, or further processed.  $\beta$ -site A $\beta$ PP cleaving enzyme-1 (BACE1) appears to act on A $\beta$ PP in late Golgi/TGN and endosomes, as indicated by the acidic optimal pH of BACE1.  $\gamma$ -secretase complex activity on the other hand takes place in multiple cellular compartments including ER, Golgi, and plasma membrane; the last is thought to comprise only a small fraction of the  $\gamma$ -secretase activity.

A key question that has existed since the elucidation of  $A\beta PP$  is the normal cellular function of this molecule, which is unresolved. One candidate ligand, secreted neuronal protein F-spondin, is implicated in neuronal sprouting and development. F-spondin binds  $A\beta PP$  as well as APLP-1 and APLP-2, which may interfere with  $\beta$ -secretase cleavage and cell signaling effected by the cytoplasmic domain (Wilquet and De Strooper 2004). A $\beta$ PP has been suggested to serve as a receptor for intracellular transport of synaptic vesicles through interaction with kinesin and microtubules (Kamal et al. 2001). Both  $A\beta$ PP and the low-density lipoprotein receptor-related protein have been shown to bind the adaptor protein Fe65 via their cytoplasmic domains which increases  $A\beta$ PP proteolytic processing (Pietrzik et al. 2004). Interestingly, both LRP and  $A\beta$ PP are also  $\gamma$ -secretase substrates after cleavage and removal of their extracellular domains. A role of  $A\beta$ PP in heavy metal binding and as an antioxidant may also be an important role with direct implications in the disease process when dysfunctional (Cho et al. 2010).

In brief, the fundamentally toxic A $\beta$ 42, otherwise a product of normal cellular metabolism, is thought to be overproduced in disease resulting in neurodegeneration, or so the amyloid cascade theory postulates. Support for this comes principally from Mendelian diseases with pathogenic A $\beta$ PP mutations leading to extensive A $\beta$  deposits and early-onset disease. In vitro toxicity of A $\beta$ 42 peptides is considered further evidence for the cascade, although toxicity lies within narrow conditions, irrelevant to brain physiology in terms of toxicity. Despite the commonly held notions, whether or not A $\beta$ 42 is toxic in vivo remains to be elucidated. On the other hand, a role of A $\beta$ 42 in neuroprotection, which is made intuitively difficult by prevailing ideas, has been demonstrated (Nunomura et al. 2006). The significance of

this cannot be understated, given again that the overwhelming majority of clinical trials accept A $\beta$ 42 toxicity as fact. The first major and somewhat infamous phase II active immunization approach (AN-1792) may have been the most informative of all trials to date given the now long-term follow-up. The evidence is now clear that removal of A $\beta$  from the brain in mild to moderate AD has no major cognitive benefits. Moreover, the two individuals studied who had almost complete removal of A $\beta$  plaques progressed to dementia at the same rate as placebo, and each expiring with a mini-mental status score of 0. This effectively answered the question of whether dementia progresses in the face of A $\beta$  removal from the brain whether it be fibrillar amyloid or oligomers (Holmes et al. 2008) and objectively speaking leaves considerable doubt as to whether A $\beta$ 42 is toxic at all. Yet the enthusiasm for anti-A $\beta$ 42 therapy is seemingly undaunted and is now progressing toward earlier intervention, based on perhaps an equally flawed notion that failure to reverse mild to moderate dementia means that intervention was too late, something for which there is no evidence whatsoever.

#### 2.1 Does Aβ Correlate with Clinical and Anatomic Indices of Disease?

While the above question data seems to have closed the case on whether A $\beta$  causes dementia (the lag in acceptance of this evidence notwithstanding), the relationship between A $\beta$  pathology and disease has been the subject of a number of studies prior to the AN-1792 trial, and indeed the data indicate unambiguously that the correlation between AB and clinical disease is imprecise at best (Castellani and Smith 2011). An early study in the 1960s showed an overall tendency toward increased disease severity with plaque burden (Blessed et al. 1968; Giannakopoulos et al. 2003), although numerous subsequent studies have refuted this concept (Giannakopoulos et al. 2003; Braak and Braak 1991; Arnold et al. 1991). At present, it is accepted that amyloid burden overall correlates poorly with disease severity, and the distribution of A $\beta$  tends to be diffused throughout the neocortex with no meaningful region specificity. Diffuse deposits of amyloid also occur in the striatum and cerebellar cortex late in disease (Montine et al. 2012), with no discernible selectivity in terms of loss of function subserved by these regions. Relative to neocortical A $\beta$ , it is of some interest that medial temporal allocortical tissue involved in memory processing shows decreased A $\beta$  (Arnold et al. 1991), something that gets little attention in the literature. Given the role of ApoE in facilitating fibrillogenesis of A $\beta$ , it is also interesting that the extent of neocortical A $\beta$  deposits shows variable correlation in the literature with ApoE susceptibility alleles, including ɛ4 (Berg et al. 1998; Nunomura et al. 2001).

A relatively new paradigm that is more functional than structural has emerged, in part, in response to the growing realization of the imprecise relationship between  $A\beta$  and clinical disease and to the clinical trials that have effectively removed  $A\beta$  and have not altered the neurodegenerative process. It is now suggested that soluble low-n  $A\beta$  oligomers cause synaptic damage and functional neurologic deficits (Selkoe 2008).

Experimental studies involving injection of conditioned medium, derived from oligomer-secreting APP V717F Chinese hamster ovary cells, into rat lateral ventricle, demonstrated alterations in long-term potentiation (LTP) that was related to low-n oligomers per se and not monomers (Walsh et al. 2002). LTP alteration was also shown in vitro in hippocampal mouse slices, along with concomitant changes in cell cycle signaling cascades and behavioral abnormalities. Here again, however, there is a premature juxtaposition of nakedly artificial experimental data and human brain function, separated from each other by so many degrees of relevance that irrelevancy is the only logical conclusion. This is in addition to lack of insight into those soluble species that are the most toxic and to the overall lack of reproducibility of many of the studies (Benilova et al. 2012). The implication is nevertheless present that these soluble low-n species, which are in vitro elaborations that cannot be directly measured or even reproduced between laboratories, are attacking the synapse, which is also not directly assessed either clinically or pathologically, and, moreover, that this combination is the true substrate for cognitive decline, even with implications for so-called mild cognitive impairment. Again, with these data in mind, the failure of treatment efforts is expected and leaves us in urgent need of a reorganization of the way AD pathogenesis is viewed by the medical and scientific communities.

### 3 Phosphorylated Tau

When the subject of the findings by Alzheimer is discussed by way of introduction to pathogenesis, it is often stated that amyloid plaques were a key discovery. The fact of the matter is that plaques were discovered some 15 years prior to Alzheimer's description, which he knew at the time, and in fact plaques were a known accompaniment of senile dementia. The first description of the neurofibrillary tangle (NFT), however, can be attributed to Alzheimer (Castellani and Smith 2011; Alzheimer 1907). It is also interesting to note that Alzheimer devoted ten sentences and two paragraphs to his initial description of the NFT, compared to only two sentences to the senile plaque (Alzheimer 1907; Wilkins and Brody 1969), suggesting that the NFT was the more intriguing, or at least novel, lesion for him. Alzheimer and his contemporaries were nevertheless hesitant to draw conclusions regarding disease pathogenesis and were more consumed with the age of onset and clinical signs which differed from "senile dementia," a known condition at that time (Moller and Graeber 1998).

Neurofibrillary tangles received relatively little attention in subsequent years, until the advent of electron microscopy (Terry et al. 1964) and, more importantly, the advent of modern molecular techniques which allowed NFT purification and identification of tau protein as a major protein component (Grundke-Iqbal et al. 1986). The enthusiasm for tau phosphorylation as a primary process in AD, however, has always been blunted by the absence of genetic linkage and the association of TAU mutations with the frontotemporal dementia clinicopathological phenotype, rather than the AD phenotype (Cairns et al. 2007). For this reason,

more than any other, phospho-tau is regarded as downstream to  $A\beta$  in terms of the pathogenic process, despite greater fidelity in its association with clinical disease (see below).

Tau is the major protein component of neurofibrillary pathology. Similar to the situation with A $\beta$ , knowledge of tau has expanded considerably since it was purified and molecular species elucidated from the insoluble lesions. We now know that the phosphorylated tau is a major protein component of neurofibrillary pathology, which in turn promoted the study of tau in copious detail.

The tau gene is comprised of over 100 kb and contains 16 exons (Hernandez and Avila 2007). Upstream of the first exon are consensus binding sites for transcription factors. Alternative splicing of tau nuclear RNA in the adult brain involving exons two, three, and ten results in six tau isoforms. These six isoforms in turn differ in the presence of either three of four repeats of 31 or 32 peptide residues in the C-terminal region (exon 10). This peptide repeat region also comprises the microtubule-binding domain and therefore has direct implications in tau pathophysiology. Moreover, tau isoforms differ in the expression of zero, one, or two inserts encoded on exons two and three. The relative amounts of these tau isoforms as well as their phosphorylation status change during development; 3-repeat tau with no inserts is expressed in the fetus and early postnatal infant, while heterogeneous isoforms are expressed in the adult brain. This switch in RNA splicing also corresponds to an overall reduction in tau phosphorylation. Tau is relatively abundant in neurons but is present in all nucleated cells given. Its major physiologic function appears to be in binding microtubules and in stabilizing microtubule assembly for polymerization.

In disease, tau is abnormally hyperphosphorylated at proline-directed serine/ threonine phosphorylation sites, including Ser-202/Thr-205 (AT8 site), Ser-214 and/or Ser-212 (AT100 site), Thr-231 and/or Ser-235 (TG3 site), and Ser-396/ Ser-404 (PHF-1 site). In addition, alternative tau splicing differs according to pathological phenotype, such that tau accumulation in AD is a mixture of 3R and 4R tau, Pick disease tends to be 3R tau, corticobasal degeneration and progressive supranuclear palsy tend to be 4R tau, and so-called argyrophilic grain disease accumulates small inclusions comprised of 3R tau.

Does phospho-tau correlate with clinical and anatomic indices of disease? In spite of the fact that tau tends to appear in the cortex subsequent to  $A\beta$  and is generally considered a secondary or downstream phenomenon (Nelson et al. 2009), it is interesting that neurofibrillary pathology correlates closely clinical signs (Braak and Braak 1991) and much more closely than  $A\beta$  deposits. Phosphorylated tau deposition, for example, has a striking tendency to involve memory circuitry early in disease as well as in the aging process (Giannakopoulos et al. 2003; Arnold et al. 1991). It is also remarkable that abundant neocortical neurofibrillary pathology is virtually always associated with clinical signs of AD (Nelson et al. 2009, 2012), whereas extensive neocortical  $A\beta$  deposits are often seen in aged individuals in the absence of significant cognitive impairment or evidence of neuronal loss. In other words, heavy tau "burden" is generally incompatible with preserved cerebral function, while heavy amyloid burden often is not. The role of phosphorylated tau in functional disease is progressing in a manner very much similar to  $A\beta$ . Recent studies, for example, indicate that insoluble tau accumulation is somewhat benign, while oligomeric phospho-tau intermediates may be more toxic and may be toxic at the specific level of the synapse (Santacruz et al. 2005). Also similar to  $A\beta$  studies, support for this concept is limited to highly experimental models (Stokin et al. 2005; Yoshiyama et al. 2007). The results highlight the growing theme that insoluble pathological lesions, and in this case NFT formation is a late stage nontoxic event, and that attention might be better directed toward upstream soluble tau intermediates. That the entire process, from changes in synthesis of soluble species, to putative soluble assembly intermediates, to insoluble pathological lesions, has yet to be embraced as a response to an underlying pathogenic process is remarkable. With recent multiple failures of the cascade concept, testing of biological alternatives is long overdue.

#### 4 Conclusion

Current hypotheses of AD pathogenesis encompass copious and sophisticated data, but nevertheless have their origin in hallmark pathological lesions described more than a century ago. The literature is now contradictory on whether hallmark lesions are best considered manifestations of neurotoxicity or instead insoluble epiphenomena to the more important events involving soluble, toxic intermediates and the synapse, events ironically that cannot be directly observed, and are based on in vitro elaborations which have proven challenging to reproduce between laboratories and have not passed the test of relevance to the human brain. More likely, and based on now considerable evidence, pathological lesions as well as their constituent proteins of whatever species, be they monomeric, oligomeric, or insoluble fibrils, can be tied to molecular pathogenesis on the basis of disease expression, or a host response. This response is likely fundamentally adaptive over a long period of time and more in line with Darwinian theory. Targeting of such lesions, or any individual protein as part of a putative pathogenic cascade for that matter, should be entertained only with considerable care, if not the sober realization that it will more likely do more harm than good, as has been demonstrated in abundance. As the pathogenesis of AD continues to be elucidated, a broadening of the focus, rather than the perseveration on a very narrow set of principles, appears to be needed.

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# **Potential Therapeutic Effects of Statins in Alzheimer's Disease**

# Cesare Mancuso, Elizabeth Head, Eugenio Barone, Marzia Perluigi, Paolo Preziosi, and D. Allan Butterfield

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C. Mancuso (🖂) • P. Preziosi

E. Head

Department of Molecular and Biomedical Pharmacology, and Sanders-Brown Center on Aging, University of Kentucky, Lexington, KY, USA

E. Barone

Institute of Pharmacology, Catholic University School of Medicine, Rome, Italy

Department of Chemistry, Center of Membrane Sciences, and Sanders-Brown Center on Aging, University of Kentucky, Lexington, KY, USA

M. Perluigi

Department of Biochemical Sciences, Faculty of Pharmacy and Medicine, Sapienza University of Rome, Rome, Italy

D.A. Butterfield

Department of Chemistry, Center of Membrane Sciences, and Sanders-Brown Center on Aging, University of Kentucky, Lexington, KY, USA

Institute of Pharmacology, Catholic University School of Medicine, Rome, Italy e-mail: cmancuso@rm.unicatt.it

#### Abstract

Statins are a family of lipid-lowering agents, long known to be beneficial in conditions where dyslipidemia occurs, such as atherosclerosis. Very recently, statins also have been proposed for use in neurodegenerative conditions, including Alzheimer's disease (AD). However, it is clear that the purported effectiveness of statins in neurodegenerative disorders is not directly related to cholesterol-lowering effects of these agents but, rather, to their pleiotropic functions.

Moreover, evidence from randomized, double-blind clinical trials demonstrated that statins have only limited beneficial effects in improving cognitive function in AD patients with moderate dementia. There is also a suggestion that in nondemented elderly people, statin use can be associated with cognitive impairments. Possible mechanisms underlying these effects are discussed along with the pros and cons of the use of statins in neurodegenerative disorders.

#### Keywords

Alzheimer's disease • Cholesterol • Dementia • Statins

#### 1 Basic Pharmacology of Statins

Statins are a family of drugs with pleiotropic functions. To this class belong eight drugs: mevastatin, lovastatin, pravastatin, and simvastatin, which are natural compounds derived from fungal products, whereas fluvastatin, atorvastatin, rosuvastatin, and pitavastatin are distinct synthetic compounds (Shitara and Sugiyama 2006). In addition, lovastatin and simvastatin have a lactone ring in the chemical structure which increases their liposolubility, whereas the remainder statins have open acid forms (Shitara and Sugiyama 2006). These differences in structure could modify the ability of individual statins to cross the blood-brain barrier.

From a pharmacodynamic viewpoint, statins inhibit hydroxy-methyl-glutaryl-CoA (HMG-CoA) reductase, which makes statins effective for the treatment of dyslipidemias (Shitara and Sugiyama 2006). By inhibiting HMG-CoA reductase, statins block the first step in cholesterol biosynthesis, namely, the conversion of HMG-CoA into mevalonate (Shitara and Sugiyama 2006; Bersot 2011). As a result of statin administration, low-density lipoprotein (LDL) cholesterol synthesis decreases in hepatocytes that in turn leads to reduced cholesterol blood level. In addition to this effect, statins are cardiovascular agents, due to their ability to counteract hyperlipidemias that are a major cause of atherosclerosis which, in turn, is a common pathogenic mechanism for coronary artery disease, ischemic cerebrovascular disease, and peripheral vascular disease (Shitara and Sugiyama 2006; Bersot 2011). In addition to this main pharmacological effect, statins are

endowed with other pleiotropic activities. Statins reduce primary cardiac risk in many types of cardiac and vascular surgeries, with greater benefits in high-risk patients (Gajendragadkar et al. 2009). Furthermore, due to the well-known immunomodulatory effect, statins prevent graft versus host disease and allogeneic transplantation (Shimabukuro-Vornhagen et al. 2009).

Although all statins share the same main mechanism of action, their pharmacokinetic profile is quite different. Statins are well absorbed by the intestine when given by oral route, even though they undergo marked first-pass effects by the liver, which reduces the systemic bioavailability (5-30%) (Bellosta et al. 2004). With the exception of simvastatin and lovastatin, which are prodrugs and require hepatic activation, other statins are administered as  $\beta$ -hydroxy-acids. Upon administration, statins reach C<sub>max</sub> (peak plasma concentration) ranging from 10 ng/ml (lovastatin and simvastatin) to 448 ng/ml (fluvastatin), with a  $T_{max}$  (time to reach  $C_{max}$ ) from 0.5–2 h (fluvastatin and pravastatin) to 2–4 h (atorvastatin, lovastatin, rosuvastatin, simvastatin) (Bellosta et al. 2004). In the bloodstream, statins are bound to albumin (43–99%), which accounts for their variable half-life. Atorvastatin and rosuvastatin are the stating with the longest half-life (15-30 h and 20.8 h, respectively), whereas fluvastatin, lovastatin, pravastatin, and simvastatin have half-lives around 0.5-3 h (Bellosta et al. 2004). All stating are metabolized by the liver through the isoforms 3A4 (atorvastatin, lovastatin, and simvastatin) and 2C9 (fluvastatin and rosuvastatin) of the cytochrome P-450 (CYP) system, with the exception of pravastatin which undergoes sulfation (Bellosta et al. 2004). The primary route of elimination is fecal, and only a minor fraction of statins is eliminated via urine (Bellosta et al. 2004; Shitara and Sugiyama 2006; Bersot 2011).

#### 2 Statins' Toxicology

The main adverse effects of statins are hepatotoxicity and myopathy. A transient elevation of serum transaminases (up to three times the baseline value) is a common outcome of statin therapy (Bersot 2011). However, the incidence of this side effect is low and dose dependent and does not imply the contraindication of statins in individuals with concomitant liver diseases such as hepatitis C (Bersot 2011). Myalgia is often associated with statin use and is paralleled by a 10-fold increase in plasma creatine kinase (Bersot 2011). Rhabdomyolysis is quite rare, and the risk of developing this side effect is correlated with the statin dose and plasma concentration (Bersot 2011). About 30 cases of serious hepatic failure and 42 cases of death due to rhabdomyolysis associated with statin administration were reported to the FDA over the last 15 years (Bersot 2011; Law and Rudnicka 2006). Particularly serious was the occurrence of fatal rhabdomyolysis in 31 patients treated with cerivastatin in the USA; among these patients, 12 received cerivastatin and gemfibrozil, thus suggesting that the cerivastatin-induced miotoxicity was potentiated by gemfibrozil (Staffa et al. 2002; Shitara and Sugiyama 2006). Due to this severe adverse effect, Bayer voluntarily withdrew cerivastatin from the market.

Rare adverse effects due to statin therapy are tendinous disorders. Tendinitis and tendon rupture were mainly associated with atorvastatin, simvastatin, and pravastatin therapy (Marie et al. 2008).

The main determinants of statin toxicity are the following.

#### 2.1 Transporters

The organic anion-transporting polypeptide (OATP1B1) is expressed in the sinusoidal membranes of hepatocytes and is the most important influx transporter of statins (Niemi 2010; Neuvonen 2010). The activity of OATP1B1, whose encoding gene SLCO1B1 undergoes genetic variability among individuals, accounts for important differences in statin plasma levels and toxicity. Two common single nucleotide polymorphism (SNP) variants of the SLCO1B1 gene were identified: c.388A>G (p.Asn130Asp; rs2306283) and c.521T>C (p.Val174Ala; rs4149056). Individuals who carry the c.388A>G (haplotype SLCO1B1\*1B) have an increased OATP1B1 activity which is associated with a 35 % reduction in pravastatin bioavailability (measured as area under the concentration-time curve, AUC) but without significant clinical outcomes (Niemi 2010). Conversely, subjects harboring the haplotypes SLCO1B1\*5 (c.521T>C) and SLCO1B1\*15 (c.521T>C together with c.388A>G) have a reduced activity of the OATP1B1 transporter and an increased statin plasma concentration (Niemi 2010). Both the latter haplotypes are differentially distributed in the population, with a combined allele frequency of 15–20 % in Caucasians, 10–15 % in Asians, and 2 % in sub-Saharan Africans and African Americans (Niemi 2010). Individuals with the c.521CC genotype have an increased bioavailability of simvastatin, pitavastatin, atorvastatin, pravastatin, and rosuvastatin of 221 %, 173 %, 144 %, 90 %, and 87 %, respectively, with respect to subjects carrying the c.521TT genotype (Niemi 2010; Neuvonen 2010). These genetic polymorphisms have indeed important clinical outcomes. The administration of 80 mg/day simvastatin to individuals with myocardial infarction was associated with myopathy, and the odds ratios were 4.5 per copy of the C allele and 16.9 in c.521CC as compared with the c.521TT homozygotes (Link et al. 2008; Niemi 2010). More than 60 % of these cases of myopathy were attributable to the C variant (Link et al. 2008). Similar results were obtained when patients were treated with 40 mg simvastatin/day with a relative risk of 2.6 per copy of the c.521C allele (Link et al. 2008; Niemi 2010). It is also important to emphasize the c.521T>C polymorphism could be responsible for the onset of myopathy even when other statins are administered. In order to reduce the risk of adverse effects, Caucasian adults harboring the SLCO1B1 c.521TC genotype should be treated with a half dose of simvastatin, atorvastatin, pravastatin, rosuvastatin, and pitavastatin, whereas those with the c.521CC genotype should receive a quarter dose of simvastatin, atorvastatin, and pitavastatin (Niemi 2010).

*ABCG2* encodes the ATP-binding cassette G2 (ABCG2), an efflux transporter located on the apical membranes of several cell types including intestinal epithelial cells and hepatocytes. The *ABCG2* gene undergoes genetic variability, and the more

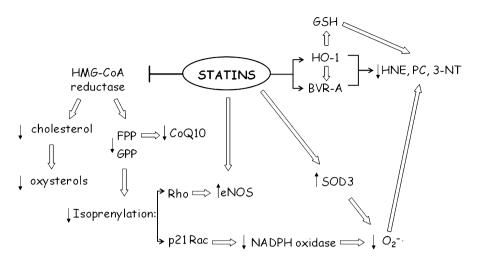
common SNP is the c.421C>A (p.Gln141Lys; rs 2231142) which reduces the transport function of ABCG2. Individuals who carry the c.421AA genotype have an increased AUC of simvastatin lactone (111 %) as well as atorvastatin and fluvastatin (72 %) with respect to those with the c.421CC genotype (Niemi 2010). Individuals with a c.421AA genotype have a greater risk to develop myopathy if treated with rosuvastatin, atorvastatin, and fluvastatin, even if no clinical trial definitely proved this possibility (Niemi 2010).

#### 2.2 Drug Interactions

In order to reduce the incidence of hepatotoxicity and myopathy, statins should not be used in association with inhibitors of CYP3A4, including grapefruit juice (see below). Several cases of rhabdomyolysis were associated with the concomitant treatment of simvastatin, lovastatin, or atorvastatin with ritonavir, cyclosporine, azole antifungals, macrolides antibiotics, and calcium channel blockers (Neuvonen et al. 2006; Neuvonen 2010). Also the combination of statins and fibrates should be avoided, in particular gemfibrozil (Bersot 2011). For further information about the potential harmful effects deriving from the interaction of statins with other drugs or nutraceuticals, see below.

## 3 Rationale for the Use of Statins in Alzheimer's Disease

Alzheimer's disease (AD) is a progressive neurodegenerative disorder characterized by cognitive dysfunction, difficulties in performing the activities of daily living, and mood changes. At autopsy, the AD brain is characterized by the accumulation of amyloid- $\beta$  peptide (A $\beta$ ) and hyperphosphorylated *tau* protein (Sultana and Butterfield 2010; Citron 2010; Querfurth and LaFerla 2010). Amyloid- $\beta$  peptide is formed by the secretase-mediated cleavage of the amyloid precursor protein (APP).  $\beta$ -Secretase cleaves APP at the N-terminus, generating an extracellular soluble fragment named sAPP $\beta$  and leaves an intramembrane fragment called C99 (Mancuso et al. 2011; Querfurth and LaFerla 2010). The latter is cleaved at the C-terminus by  $\gamma$ -secretase and A $\beta$  is released (Mancuso et al. 2011; Querfurth and LaFerla 2010). Once formed, A $\beta$  aggregates as oligomers and fibrils, which form the central core of senile plaques (Sultana et al. 2009; Citron 2010; Querfurth and LaFerla 2010). Current research suggests that A $\beta$  oligomers are more toxic than aggregates (Shankar et al. 2008; Butterfield et al. 2007; Querfurth and LaFerla 2010), and this evidence provides a plausible explanation of the reason why drugs targeted to facilitate the disaggregation of A $\beta$  fibrils into oligomers, e.g., bapineuzumab and solanezumab, failed to improve cognitive function in AD patients (Ayrolles-Torro et al. 2011; Mancuso et al. 2011). Interestingly, the gene encoding APP is located on the chromosome 21; therefore, individuals with down syndrome (DS), who have 3 copies of this chromosome, have an increased risk of developing AD (Bush and Beail 2004). Autopsy findings in individuals with DS



**Fig. 1** Cholesterol-dependent and cholesterol-independent effects of statins and repercussion on the redox status of the cell. For further information, see text. Abbreviations: *3-NT* 3-nitrotyrosine, *BVR-A* biliverdin reductase-A, *CoQ10* coenzyme Q10, *eNOS* endothelial nitric oxide synthase, *FPP* farnesyl pyrophosphate, *GPP* geranyl pyrophosphate, *GSH* reduced glutathione, *HMG-CoA* hydroxyl-methyl-glutaryl-CoA, *HNE* 4-hydroxynonenals, *HO-1* heme oxygenase-1, *PC* protein carbonyls, *SOD3* extracellular superoxide dismutase

revealed senile plaques and neurofibrillary tangles in the brain, which are secondary to the formation and deposition of A $\beta$  and tau protein, respectively (Gyure et al. 2001; Itoh and Yagishita 1998).

Amyloid- $\beta$  peptide and tau protein synergize and exert their neurotoxic effects through several mechanisms: (a) generation of superoxide anion and nitric oxide through the activation of NADPH oxidase and inducible nitric oxide synthase (iNOS), respectively (Butterfield et al. 2007; Querfurth and LaFerla 2010), which react each other and form peroxynitrite (ONOO<sup>-</sup>); (b) mitochondrial impairment secondary to the inhibition of key enzymes involved in the respiratory chain and Krebs cycle (Querfurth and LaFerla 2010); and (c) stimulation of the ionotropic glutamate receptor NMDA and increase of  $Ca^{2+}$  overload thus leading to excitotoxic cell death (Querfurth and LaFerla 2010). Taking into consideration these multiple toxic pathways, statins may be particularly useful for AD. Statins can inhibit endothelial  $O_2^{-}$  formation by preventing the isoprenylation of p21 Rac, which is critical for the assembly of NADPH oxidase after activation of protein kinase C (Wallerath et al. 2003) (Fig. 1). In addition, the cellular superoxide clearance activity is increased given that SOD3 activity is more than doubled by simvastatin. Further, simvastatin treatment also increases the number of functionally active endothelial progenitor cells (Landmesser et al. 2005). Moreover, statins increase the expression of endothelial nitric oxide synthase (eNOS) through the inhibition of Rho isoprenylation (Laufs et al. 1998), and statins can also directly activate eNOS via posttranslational mechanisms involving activation of the PI3K/Akt pathway (Kureishi et al. 2000). Statins showed positive effects against A $\beta$ -induced oxidative stress in mouse models of AD (Kurinami et al. 2008; Tong et al. 2009) as well as a reduction in the phosphorylation of tau protein in human cerebrospinal fluid (CSF) (Riekse et al. 2006). However, although statins' treatment appears to provide benefits, it is difficult to determine whether the benefits are due to lower cholesterol levels or to statin pleiotropy. Atorvastatin treatment was neuroprotective against cell degeneration induced by A $\beta$ (1-40), reducing inflammatory and oxidative responses and increasing the expression of glutamatergic transporters (Piermartiri et al. 2010). Murphy et al. showed that longterm atorvastatin did not reduce  $A\beta$  levels, despite a significant reduction in  $\beta$ -secretase 1 (BACE1) protein levels and activity in the brains of aged beagles (Murphy et al. 2010). Subsequently, Barone et al. found that although no changes in Aß levels occur, long-term atorvastatin significantly reduced lipid peroxidation, protein oxidation and nitration, and increased GSH levels in the parietal cortex of treated dogs (Barone et al. 2011). This effect was cholesterol and AB independent and specific for brain (Barone et al. 2011). Another possible mechanism by which statins exert neuroprotective effects is the enhancement of the cell stress response. The administration of atorvastatin 80 mg/day for 14.5 months to aged beagles increased the expression of the heme oxygenase-1/biliverdin reductase-A (HO-1/ BVR-A) system, a main effector of the adaptive stress response (Mancuso and Barone 2009), and this correlated with a significant reduction in both oxidative and nitrosative stress biomarkers in the parietal cortex of treated beagles that showed improved cognition (Barone et al. 2012; Butterfield et al. 2011). Conversely, side effects of long-term statin treatment include a decrease in serum and tissue CoQ10 levels, which could result in an energy metabolism impairment in heart, skeletal muscle, and liver (Martin et al. 2011; Bliznakov and Wilkins 1998). The importance of statin-induced energy metabolism impairment and its role in the development of their side effects was demonstrated also in humans. The administration of 240 mg/day CoQ10 reverted fatigue, myalgia, and peripheral neuropathy in 50 patients chronically treated with statins (Langsjoen et al. 2005).

More detailed research into the pharmacology of statins is necessary, particularly the concentrations achieved in the central nervous system and the level at which they block the production of cholesterol and they modulate all the above pathways, in order to prove their beneficial effect and support the potential use of statins in neurodegenerative disorders.

### 4 Statins and Dementia: The Support Given by the Evidence-Based Medicine

Although the majority of the preclinical evidence shows neuroprotective effects of statins in ameliorating cognitive dysfunction, clinical data largely do not show similar benefits. In addition, some clinical studies show the opposite results depending on whether they are observational studies or randomized clinical trials (RCT).

Early cohort and case-control studies demonstrated that statins reduced the risk of developing dementia, including AD, and this protective effect was maintained over a 6-year follow-up period (Beydoun et al. 2010; Li et al. 2010; Smeeth et al. 2009; Haag et al. 2009; Horsdal et al. 2009; Rosenberg and Allard 2008; Sparks et al. 2008; Cramer et al. 2008). These findings were recently challenged by Benito-Leon et al., who demonstrated that statins did not improve cognitive performance in nondemented elderly subjects with a median age of 72 years (Benito-Leon et al. 2010). Similar results were obtained in a large cohort study, which involved more than two million participants aged 30–84 years of whom 10.7 % received statins (Hippisley-Cox and Coupland 2010). A cross-sectional research study, with a follow-up of 5 years, demonstrated that in 123 Caucasian subjects with DS (41–78 years), total cholesterol levels > 200 mg/dl positively correlated with the development of AD (hazard rate 2.59), whereas individuals taking statins had less than half the risk of developing AD with respect to nonusers (hazard rate 0.402) (Zigman et al. 2007).

In order to confirm these epidemiological studies, some RCT were performed with comparable results. The PROSPER study, which enrolled approximately 6,000 people aged 70-82 years, demonstrated that pravastatin (40 mg/day) did not have any effect on cognitive function over a follow-up of 3 years (Trompet et al. 2010; Shepherd et al. 1999). The LEADe study tested the hypothesis that atorvastatin (80 mg/day) over 72 weeks delayed cognitive decline in patients with mild-tomoderate AD. The results of this study did not support any significant positive effect of atorvastatin on cognitive or global function in patients receiving the statin compared to those with placebo (Feldman et al. 2010). In August 2011, Sano et al. published the results of an RCT of simvastatin (20 mg/day for 6 weeks and 40 mg/day for the remainder 18 months) in participants with mild-to-moderate AD that also demonstrated the lack of efficacy (Sano et al. 2011). On the other hand, the ADCLT trial demonstrated that atorvastatin (80 mg/day) for 1 year exhibited a significant positive effect on cognitive performance after 6 months of therapy compared with placebo. However, this beneficial effect was selective for individuals who matched restricted criteria, such as a higher MMSE score at baseline, total cholesterol levels higher than 200 mg/dl, and the presence of an apolipoprotein E-4 allele (APOE-4) (Sparks et al. 2006a, b). When all the RCT study results are combined, a meta-analysis suggests that there is insufficient evidence to recommend statins for the treatment of dementia and AD (McGuinness et al. 2010). This statement matches the guidelines of the British Association for Psychopharmacology who also do not recommend statins for the prevention or treatment of AD (O'Brien and Burns 2010). However, in a pilot clinical trial, a modest improvement in cognition was detected in individuals with mild cognitive impairment (MCI), arguably the earliest form of AD, treated with statins (Sparks et al. 2010).

### 5 Unresolved Issues

In order to understand the reasons for the inconsistent beneficial effects of statins in aged or demented individuals, several issues need to be addressed and carefully evaluated.

Ability to Cross the Blood-Brain Barrier. As mentioned earlier, the presence of a lactone ring in the chemical structures of simvastatin and lovastatin makes these latter very liposoluble, whereas the open acids atorvastatin, fluvastatin, and pitavastatin have an intermediate liposolubility, and pravastatin has the lowest one (Shitara and Sugiyama 2006). Such different degree of liposolubility might suggest the use of lipophilic statins in demented patients to increase the bioavailability in the brain. However, both the LEADe and CLASP trials failed to demonstrate a beneficial effect of lipophilic atorvastatin and simvastatin on cognitive function in AD patients, similar to the results obtained in the PROSPER study involving the lipophobic pravastatin. Thus, the different degrees of liposolubility are unlikely to be a key determinant of the limited effectiveness of statins in these clinical trials.

Age. Epidemiological data demonstrate that the incidence of AD increases with age and doubles every 5 years after 65 years of age with 1,275 new cases/1,00,000 persons/year (Querfurth and LaFerla 2010). In the Western hemisphere, the prevalence of AD was calculated at approximately 1% in subjects aged 60-64 but increases to between 33 % and 50 % in people aged 85 or older (Mayeux 2010). Much of the epidemiological and clinical studies designed to examine the role of statins in AD enrolled included individuals aged 65–84 years. Although the clinical studies and the meta-analysis discussed above did not support an overall beneficial effect for statins in dementia and AD, it is noteworthy to mention that those studies in which stating had a major effect on cognitive functions included individuals aged 68-74 years (Simons et al. 2002; Li et al. 2010; Haag et al. 2009). Consequently, it is possible that 68-74 years of age should be considered an optimal age for statin efficacy in preventing dementia. On the other hand, both the LEADe and PROSPER studies, which recruited individuals within the same range of age, failed to demonstrate any beneficial effect of atorvastatin and pravastatin in people with AD. Subjects aged 80 or older also did not have any beneficial effects from statins (Li et al. 2010).

Cholesterol Blood Levels Before and After Treatment with Statins. An important question to address when studying the HMG-CoA reductase-independent effects of statins is the concomitant reduction of LDL cholesterol plasma levels. This issue is quite important when the pleiotropic effects of statins are studied in the nervous system because cholesterol is a main component of cell membranes, in particular myelin (Saher et al. 2005), and if cholesterol blood levels fall due to uncontrolled therapy with lipid lowering agents, nervous function would also decrease. The majority of AD patients recruited for clinical trials had serum LDL cholesterol at baseline around 131–147 mg/dL, which was significantly reduced by 50–54 % after the administration of atorvastatin (80 mg for 52 weeks) or simvastatin (40 mg for 26 weeks) (McGuinness et al. 2009; McGuinness et al. 2010). These values of LDL cholesterol, before and after statin treatment, are acceptable and do not imply any possible adverse effects. However, although there is a marked effect of statins on LDL cholesterol plasma levels, no beneficial effects on cognitive function are observed in normocholesterolemic patients (McGuinness et al. 2009; McGuinness et al. 2010). Indeed, even in hypercholesterolemic patients, the administration of statins did not have any disease-modifying effect on AD (Hoglund et al. 2004;

Riekse et al. 2006). Evans et al. showed that in AD patients heterozygous for APOE-4 allele or carriers of PS1 mutations, the administration of simvastatin or atorvastatin only slightly reduced the concentration of CSF cholesterol at 6–7 months followed by a peak at 2 years and a return to baseline levels after 3 years (Evans et al. 2009). This finding suggests that despite changes in plasma cholesterol levels, statins cause only minimal changes in brain cholesterol, and, therefore, the effect on cognitive functions may be independent of systemic cholesterol metabolism.

Interaction with Xenobiotics. Patients with AD, as well as other types of dementia, usually take additional drugs for other age-related disorders or comorbidities. As mentioned above, all statins, with the exception of pravastatin, are metabolized by CYP3A4 or CYP2C9, and their plasma levels could be reduced or increased with concomitant administration of drugs that induce or inhibit these CYP isoforms. Donepezil and galantamine, two acetylcholinesterase inhibitors provided to AD patients, are metabolized by CYP3A4 (Jann et al. 2002) and, therefore, could compete with statins. As a consequence of this competition, statin plasma levels could increase as well as the risk of side effects. Also, AD patients may also be supplemented with nutraceuticals including curcumin, green tea extracts, or grapefruit juice as these natural substances are widely considered to be free radical scavengers and therefore neuroprotective. Unfortunately, these natural substances are inhibitors of CYP3A4 and, therefore, increase plasma concentrations of statins (Kiani and Imam 2007; Stump et al. 2006; Hare and Elliott 2003). Interestingly, it was reported that the concomitant administration of simvastatin and consumption of grapefruit or green tea causes rhabdomyolysis (Dreier and Endres 2004; Werba et al. 2008).

Taking into consideration the results from evidence-based medicine, it is possible to suggest that select individuals with AD may benefit more from statins: aged 65–74 years, carrier of an ApoE-4 allele (Sparks et al. 2006a) but without the haplotype *SLCO1B1\*5* or *SLCO1B1\*15* or *SLCO1B1\*1B/\*1B* (Niemi 2010), normocholesterolemic and sparing user of drugs which inhibit CYP3A4.

#### 6 Conclusion

Although preclinical data suggest beneficial effects of statins in the treatment of dementia and AD, results derived from epidemiological studies and RCT are contradictory. Several critical issues may need to be considered (1) the genetic profile of individuals with regard to specific genes involved in statin absorption, (2) the rate of inhibition of cholesterol synthesis in normocholesterolemic patients, (3) age of the individual, and (4) concomitant administration of drugs or nutraceuticals. Thus, future RCT using statins may consider from recruiting AD patients aged 65–75 years, which is an age range that appears to benefit from statin therapy, with primary endpoint, the analysis of cognitive function, and a significant longer follow-up. Another suggestion is to study the effect of statins in individuals with MCI in order to understand whether or not the neuroprotective effect of statins could block or slow the transition to AD. This recommendation could be especially

important since preclinical studies with atorvastatin decreased oxidative stress in aged dogs, and this benefit was correlated with levels of A $\beta$ (1–42), which is same sequence as that of humans (Barone et al. 2011). Oxidative stress is strongly associated with amnestic MCI and AD (Perluigi et al. 2009; Sultana et al. 2010; Sultana et al. 2009), and A $\beta$  is hypothesized to contribute to this oxidative stress (Butterfield et al. 2001).

In conclusion, the current clinical evidence is not strong enough to support the widespread use of statins to treat dementia and AD. However, it is critical that researchers and clinicians in the near future investigate whether or not statin therapy should be restricted to selected populations of demented individuals with the best chance of efficacy derived from evidence-based medicine is recommended.

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